<u>Homework 3 (100 points)</u> (Regression & Classification)

This assignment has 3 separate problems. Each problem will require you to create a separate source code pipeline (using a jupyter notebook) and perform all the requisite analyses. In addition to performing the analysis (i.e., writing code in the jupyter notebook), any questions that have been posed in the word document can be answered directly inline (i.e., below each question in the document itself). The assignment is worth 100 points in total.

What to submit:

- 1. For each question, you will be provided with a separate jupyter notebook to give your requisite code / analysis.
- In addition, you can use this word document to answer all the non-coding questions (inline) and submit a single document for all the analysis.

<u>Submission format:</u> Do not modify the directory structure of the zip archive that you download to start the assignment. Essentially, make sure each jupyter notebook is in a separate directory (which is how the assignment is initially provided i.e., with Q1, Q2, Q3 subdirectories). The *HW3_Questions_and_solutions.pdf* (i.e., the current file with the solutions that you will fill in and convert to PDF format), should be placed at the root of the zip archive.

Once all the solution pdf and all the source code is complete, zip the full assignment and submit the zip archive. Your submission zip archive should have the following structure:

<lastname_firstname_CS556_HW3.zip>/ ./HW3_Questions_and_Solutions.pdf Q1/Q1.ipynb Q2/Q2.ipynb Q3/Q3.ipynb

Leave the datasets unchanged and in the directory itself. Each of your jupyter notebooks should be directly executable without any further modifications (i.e., if your assignment submission is downloaded from canvas and unzipped), any of your submitted notebooks if executed using a jupyter server should be directly runnable without additional data downloads etc.

Q1. Housing Price Prediction (34 points)

a. **Dataset Description:** The data pertains to the houses found in each California district and some summary statistics about them based on the 1990 census data. It contains one instance per district block group. A block group is the smallest geographical unit for which the U.S. Census Bureau publishes sample data (a block group typically has a population of 600 to 3,000 people).

The goal of this task is to design a regression model to predict the *median house value* conditioned upon a set of input attributes corresponding to a particular California district block.

The attributes in the dataset are as follows; their names are self-explanatory:

- i. <u>longitude (continuous)</u>: One of the coordinates that are used to identify the California district block
- ii. <u>latitude (continuous)</u>: One of the coordinates that are used to identify the California district block
- iii. housing median age (continuous): Average age of the house in California district block
- iv. total rooms (continuous): Total number of rooms of all the houses in the California district block
- v. total bedrooms (continuous): Total number of bedrooms of all the houses in the California district block
- vi. population (continuous): Number of people residing in the district block
- vii. households (continuous): Number of families in the district block
- viii. median_income (continuous): Median income for households in the district block of houses (measured in tens of thousands of US Dollars)
- ix. ocean proximity (categorical): Location of the house. Is it inland, near the bay, near the ocean, etc.
- x. <u>median_house_value.(continuous):</u> Median house value within a district block (measured in US Dollars)

Our target variable will be <u>median house value</u>. Use the rest of the fields mentioned above to predict the <u>median house value</u>.

b. Data Loading / Preprocessing: (10 points)

- i. Loading: (2 points)
 - 1. Load the California housing dataset using <u>pandas.read_csv()</u> function and store it in the variable (i.e., a pandas dataframe) named 'df'.
 - 2. The resulting data frame should have the shape (20942, 10) indicating that there are 20942 rows and 10 columns.
 - 3. Find the missing values in the data frame. If any (i.e., even if one column in each instance / row has a missing value), drop the row using <u>pandas.DataFrame.dropna()</u> function. The resulting data frame should have the shape (20735, 10) indicating that there are 20735 rows and 10 columns.
 - 4. Now plot a box plot for median house values to check for extreme values using df.boxplot() function.
 - 5. Now filter the dataframe and remove all the values that are above 1.5iqr. You can find the iqr by subtracting the 75th and 25th quantile using the quantile function in pandas. You can then find the upper and lower limit by adding and subtracting 1.5*iqr to the 75th quantile and 25th quantile respectively.
 - 6. You should see that the new shape of the dataframe is (19557,10).

7. Create a data frame 'corr_df' by dropping the columns *latitude*, *longitude*, and *ocean_proximity* using the <u>pandas.DataFrame.drop()</u> function. Use the *Pearson correlation* to find the correlation of each remaining feature in the 'corr_df' with the target variable 'median_house_value' using the function pandas.DataFrame.corrwith(). Report results in the following table.mod

| Feature Name | Pearson Correlation |
|--------------------|---------------------|
| housing median age | 0.068098 |
| total_rooms | 0.144141 |
| total bedrooms | 0.076512 |
| population | 0.015151 |
| households | 0.095864 |
| median_income | 0.618104 |

8. Create a data frame X of features (by dropping the column 'median_house_value'' from the original data frame) using the pandas.DataFrame.drop() function. Create a Series object of targets Y (by only considering the 'median_house_value' column from the original data frame (Do NOT use the 'corr_df' data frame in this step. Use the data frame which was obtained as a result of step b.i.3 above).

ii. Data Visualization: (3 points)

- 1. Use <u>pandas.DataFrame.hist(bins = 50)</u> function for visualizing the variation on the columns housing_median_age, total_rooms, total_bedrooms, population, household, median_income and median_house_value. Plot each histogram as a separate subplot.
- 2. Use <u>pandas.dataframe.describe()</u> function to find the mean, median and standard deviations for each feature and report in the jupyter notebook.
- 3. Use <u>pandas.get_dummies</u> to convert categorical variables into dummy /one-hot encoding. In this case the categorical column is *ocean_proximity*

iii. Data Splitting: (2 points)

Split data into training and test sets using the sklearn <u>train test split()</u> function. Perform 70-30 distribution i.e. 70% training and 30% testing. The result of your data split should yield 4 separate data frames X_train, X_test, y_train, y_test. (respectively, the training features, testing features, training targets and testing target).

iv. Data Scaling: (3 points)

- Use the <u>StandardScaler()</u> to instantiate the standard scaler class. **Note**: You will need two separate scaler objects, one to scale the features, another to scale the target values.
- 2. For each scaler, employ the 'fit_transform()' function (only on the training features, training targets) of the scaler to retrieve the new (scaled) version of the data. Store them in *X_train*, and *y train* again.
- Scale the X_test and y_test as well and store the scaled values back in X_test and y_test. (i.e., use
 the appropriate "fitted" scaler above to "transform" the test data. Note: the function to be employed
 in this case is 'transform()' as opposed to 'fit_transform()').
 Henceforth, X_train, y_train, X_test, y_test will refer to the scaled data unless stated otherwise.
- 4. Use <u>pandas.DataFrame.hist(bins = 50)</u> function for visualizing the variation of numerical attributes *housing_median_age, total_rooms, total_bedrooms, population, household, median_income* and *median_house_value* for the *X_train* and *y_train* dataset (similar to step b.ii.1 above). Once again, plot each histogram as a separate subplot.

c. Modelling: (10 points)

i. Employ Linear Regression from sklearn.linear model, and instantiate the model.

- ii. (3 points) Once instantiated, 'fit()' the model using the scaled X_train, y_train data.
- iii. (2 points) Employ the 'predict()' function to obtain predictions on X_test. Store the predictions in a variable named 'y_preds'. Note: Since the model has been trained on scaled data (i.e., both features and targets, the predictions will also be in the "scaled" space. We need to transform the predictions back to the original space).
- iv. (2 points) Use <u>inverse_transform()</u> function to convert the normalized data ('y_preds') to original scale. Store the transformed values back into 'y preds'.
- v. (3 points) Perform \underline{PCA} on the features (X train) and set n components as 2.
 - Show a scatter plot where on the x-axis we plot the first PCA component and second component on the y-axis.
 - 2. Calculate the total percentage of variance captured by the 2 PCA components using pca.explained variance ratio. Also, report the strength of each PCA component using pca.singular values.

d. Evaluation: (6 points)

- i. (2 points) Plot a scatter plot using <u>matplotlib.pyplot.scatter</u> function. Plot the predicted median house values on the y-axis vs the actual median house values on the x-axis.
- ii. (2 points) Calculate MAPE, RMSE and R² for the model and report them in the following table.

| Model | MAPE | RMSE | \mathbb{R}^2 |
|-------------------|--------------------|-------------------|---------------------|
| Linear Regression | 1445967.6798857003 | 205344.8003799818 | -40550723717.257416 |

iii. (2 points) To check how the model performs without removing the outliers, read the data frame again, and remove the missing values, scale it and retrain the model using sklearn and report the MAPE, RMSE and R² in the format specified in evaluation #2.

| Model | MAPE | RMSE | \mathbb{R}^2 |
|-------------------|-------------------|-------------------|--------------------|
| Linear Regression | 1803264.708641179 | 228929.2141310129 | -50837818649.29756 |

e. Discussion: (8 points)

i. (2 points) Based on the weights of the linear regression model, rank the features (note: only continuous features) in decreasing order of influence on the predictions.

Features ranked by importance:

| | Feature | Importance |
|---|--------------------|------------|
| 0 | longitude | 0.819479 |
| 1 | latitude | 0.764797 |
| 7 | median_income | 0.483271 |
| 5 | population | 0.338945 |
| 4 | total_bedrooms | 0.210746 |
| 6 | households | 0.142078 |
| 2 | housing_median_age | 0.109366 |
| 3 | total rooms | 0.026813 |

- ii. (2 points) Discuss how the influence of the features (obtained in question e.1) relates to the pair-wise correlation results calculated above i.e., are the features that are highly correlated with the output also the most influential or is there some other phenomenon being observed?
 - It seems like features might interact with each other in ways that aren't captured just by correlations. For example a feature may only become important when another feature exists.
- iii. (2 points) Comment about the MAPE, RMSE, R² results. What can we learn from each of these results about the model prediction performance?
 - i. MAPE Measures the average percentage error between the predicted and the actual values. The MAPE percent is the percent that the models' predictions are off by on average.
 - ii. RMSE Measures the square root of the squared difference between the predicted and actual values and is good for telling you how much deviation to expected in your predictions.

- iii. R² Tell us how much of the variance of the dependent variable can be predicted from the independent variable.
- iv. (2 points) Why is centering and scaling the data important before performing PCA?
 - i. Centering ensures that the first principal component doesn't just point toward the mean instead of reflecting the direction of the maximum variance.
 - ii. So that no one component has undue influence over the calculation

Q2. Titanic Classification Problem (33 points)

- a. Dataset Description: The sinking of the Titanic is one of the most infamous shipwrecks in history. On April 15, 1912, during her maiden voyage, the widely considered "unsinkable" RMS Titanic sank after colliding with an iceberg. Unfortunately, there weren't enough lifeboats for everyone onboard, resulting in the death of 1502 out of 2224 passengers and crew. While there was some element of luck involved in surviving, it seems some groups of people were more likely to survive than others. Our goal is to develop a classifier to predict whether a passenger survived the calamity. The attributes in the datasets are as follows:
 - i. PassengerID (categorical): Passenger ID
 - ii. Ticket (categorical): Ticket Number
 - iii. Name (categorical): Passenger Name
 - iv. Cabin (categorical): Passenger Cabin
 - v. Pclass (categorical): Passenger Class (1 = 1st; 2 = 2nd; 3 = 3rd)
 - vi. Sex (categorical): Passenger Sex
 - vii. Age (continuous): Passenger Age
 - viii. SibSp (continuous): Number of Siblings/Spouses Aboard
 - ix. Parch (continuous): Number of Parents/Children Aboard
 - x. Fare (continuous): Passenger Fare
 - xi. Embarked (categorical): Port of Embarkation (C = Cherbourg; Q = Queenstown; S = Southampton)
 - xii. Survived (categorical): Survival (0 = No; 1 = Yes) --- this is the target column. Make sure to encode data appropriately for each classifier.
 - xiii. Our target variable will be <u>Survived</u>. Use the rest of the fields mentioned above to predict whether a passenger survived the Titanic shipwreck.

b. Data Loading / Preprocessing (10 points)

- i. Loading (4 points)
 - Load the data <df_train.csv> and <df_test.csv> as a pandas dataframe using the <u>'pandas.read_csv'</u> function. The 'df_test.csv' has been preprocessed (I.e., null values have been dropped, certain columns etc. have been dropped) and should not be changed apart from splitting the the dataframe into X_test and y_test. The 'df_train' data has NOT been preprocessed and you will need to preprocess and prepare the 'df_train' dataframe. Note: Neither df_train nor df_test have been scaled. The next few steps will the enumerate data preprocessing, scaling requirements we need to perform.
 - 2. The resulting dataframe (i.e., *df_train*) should have the shape (712,12) indicating that there are 712 instances and 12 columns.
 - 3. In *df_train*, dataframe, currently you have 12 columns which are the following PassengerID, Pclass, Name, Sex, Age, SibSp, Parch, Ticket, Fare, Cabin, Embarked and the Survived column (target variable).
 - 4. Use the <u>'pandas.isnull()</u>.sum()' function check if there are any missing values in the df_train dataframe. Report which columns have missing (i.e., null) values and provide the number of the null values in the columns.
 - 5. Use the 'pandas.DataFrame.drop()' function to drop the 'Cabin', 'PassengerID', 'Name' and 'Ticket' columns.
 - 6. Use the 'pandas.DataFrame.fillna()' function to replace the NA values in the 'Age' column with the mean value of the 'Age' column. Note: This process is called 'imputation' (i.e., filling null values with a pre-specified value) and we are employing one strategy called *mean imputation*, but other strategies can also be employed in general. Use the 'dropna()' function to drop any remaining rows that consist of NA values.
 - 7. Your task is to use the feature columns to predict the target column. This can be cast as a classification problem.
 - 8. Create a pandas dataframe X_train of features (by dropping the 'Survival' column from the df_train dataframe). Create a pandas *Series* object of targets y_train (by only considering the 'Survival' column from the df_train dataframe). Moving forward, we will be working with X_train and y_train. At this point also split the df_test into X_test and y_test by dropping the 'Survival' column and storing the features into X_test. Store only the 'Survival' column in y_test.
- ii. Data Visualization (4 points)
 - 1. Using matplotlib employ a scatter plot using 'matplotlib.pyplot.scatter' between the age of the passengers and the price of their fare. Label the x-axis and the y-axis along with the giving the plot a title.
 - a. What is the highest and lowest fare price?

i. Max: 512.329200 ii. Min: 0.000000

b. What are the respective mean values of these two features?

i. Mean: 32.658449

- c. What was the age of the oldest passenger? 80
- 2. Only for this question use the df_train dataframe. Using matplotlib visualize the number of males and females that survived and their respective *passenger classes* on two separate bar chart plots using 'matplotlib.pyplot.bar' (Passenger Class column).
 - a. Which *class of passengers* had the least number of survived males and how many? Repeat this analysis for females.

i. Males: 2, 15

- ii. Females: 2, 50
- b. Which *class of passengers* had the greatest number of survived males and how many? Repeat this for females.
 - i. Males: 1, 45 ii. Females: 1, 75
- 3. Using the Target variable (Survived) in y_train, plot a bar chart showing the distribution of the 'Survived' column.
 - a. What initial comment can you make about this distribution in terms of the frequency of each class?
 - i. More people died than survived.
- 4. So far you should have successfully been able to load, preprocess and visualize your data. Use the 'pd.get dummies()' function to convert categorical data into dummy variables ('Sex' and 'Embarked'). Make sure to pass 'drop_first=True' to the 'get_dummies()' function. (Perform this only on X train store the result back into X train).
 - a. What is the new shape of X?
 - i. 710, 8
- iii. Data Scaling (2 points)
 - 1. Employ X_train and MinMaxScaler only on the continuous attributes. Employ the 'fit transform()' function of the scaler to retrieve the new (scaled) version of the data. Store the scaled values in X_train again.
 - Scale the X_test using the scaler you have just fit, this time using the <u>`transform()`</u> function.
 Note: store the scaled values back into X_test. At the end of this step, you must have X_train, X test, all scaled according to the MinMaxScaler.

c. Modelling (10 points)

- i. (2 points) Modelling (Model Instantiation / Training) using **Logistic Regression classifier**
 - 1. Employ the <u>Logistic Regression</u> classifier from sklearn and instantiate the model. Label/Name this model as 'model_lr'
 - 2. Once instantiated, 'fit()' the model using the *scaled* X_train, y_train data.
 - 3. Employ the `predict()` function to obtain predictions on X_test and store this in a variable named 'y_pred_lr'.
 - 4. Employ the <u>'accuracy_score'</u> function by using the 'y_pred_lr' and 'y_test' variables as the functions parameters and print the accuracy of the Logistic Regression model.
- ii. (2 points) Modelling (Model Instantiation / Training) using Support Vector Machine Classifier
 - Employ the <u>Support Vector Machine (SVM)</u> classifier from sklearn (SVC) and instantiate the model. Make sure to set 'probability=True' when creating the model. Label/Name this model as 'model svm'
 - 2. Once instantiated, 'fit()' the model using the scaled X_train, y_train data.
 - 3. Employ the 'predict()' function to obtain predictions on X_test and store this in a variable labeled as 'y_pred_svm'.
 - 4. Employ the <u>'accuracy_score'</u> function by using the 'y_pred_svm' and 'y_test' variables as the functions parameters and print the accuracy of the SVM model. and print the accuracy of the SVM model.
- iii. (3 points) Modelling Logistic Regression Classifier with the addition of noise on the target variable. In the data repository you should see three noisy datasets 'df train noise20', 'df train noise40',

'df train noise60'. These datasets have already been preprocessed. In each dataset

'df_train_noise<integer>, the integer indicates the percentage of noise injected into the target variable in that training set. The noise can be considered a result of incorrect class labelling of a particular instance. For example, in df_train_noise20, 20% of the instances have an incorrect target label in the training set. Our goal will be to train a set of classification models on such noisy training data and test on a clean test set (i.e., same as what we have been using so far 'df test').

Load the 'df_train_noise<nl>' (nl means noise level and is a place holder for the integer percentage) datasets and split the data into X_train_<nl> and y_train_<nl> (e.g., if working with `df_train_noise20' we would split the data and store it in variables named X_train_20, y_train_20). 'X_train_<nl>' should store the features and 'y train <nl>' should store the target variable.

Repeat the following steps (1-4) for the 20%, 40%, 60% noise level datasets.

Train a new Logistic Regression model on the new training and use the pre-existing X_test and y_test to evaluate your model. Label/Name this model as 'model lr noise <nl>'. Specifically, do the following:

- 1. Employ a new Logistic Regression classifier from sklearn and instantiate the model. Label this model as 'model lr noise<nl>'
- 2. Once instantiated, 'fit()' the model using the X train <nl> and y train <nl> data.
- 3. Employ the 'predict()' function to obtain predictions on X_test and store this in a variable labeled as 'y pred lr noise<nl>'.
- 4. Employ the <u>'accuracy score'</u> function and print the accuracy of the new Logistic Regression model.
 - a. What is your initial observation of the accuracy, is the accuracy higher or lower than that of the clean dataset?
 The clean dataset is substantially higher accuracy on the clean data.
- iv. (3 points) Modelling Support Vector Machine (SVM) Classifier with the addition of noise on the target variable. In the data repository you should see three noisy datasets 'df_train_noise20', 'df_train_noise40', 'df_train_noise60'. These datasets have already been preprocessed. In each dataset 'df_train_noise<integer>, the integer indicates the percentage of noise injected into the target variable in that training set. The noise can be considered a result of incorrect class labelling of a particular instance. For example, in df_train_noise20, 20% of the instances have an incorrect target label in the training set. Our goal

will be to train a set of classification models on such noisy training data and test on a clean test set (i.e.,

Load the 'df_train_noise<nl>' (nl means noise level and is a place holder for the integer percentage) datasets and split the data into X_train_<nl> and y_train_<nl> (e.g., if working with `df_train_noise20' we would split the data and store it in variables named X_train_20, y_train_20). 'X_train_<nl>' should store the features and 'y train <nl>' should store the target variable.

Repeat the following steps (1-4) for the 20%, 40%, 60% noise datasets.

same as what we have been using so far 'df test').

Train a new SVM classification model on the new training and use the pre-existing X_test and y_test to evaluate your model. Label/Name this model as 'model svm noise <nl>'. Specifically, do the following:

- 1. Employ a new SVM classifier from sklearn and instantiate the model. Label/Name this model as 'model_svm_noise<nl>'
- 2. Once instantiated, 'fit()' the model using the X_train_<nl> and y_train_<nl> data.
- 3. Employ the 'predict()' function to obtain predictions on X_test and store this in a variable labeled as 'y_pred_svm_noise<nl>'.
- Employ the <u>'accuracy_score'</u> function and print the accuracy of the new Logistic Regression model.
 - a. What is your initial observation of the accuracy, is the accuracy higher or lower than that of the clean dataset?
 - i. It is lower than that of the clean dataset

d. Evaluation (5 points)

- i. (3 points) Report F1 Score, Precision, Recall, Accuracy (All on the test set X test, y test)
 - Employ a <u>`classification_report()`</u> function from sklearn.metrics to report the precision recall and fl score for each class for the 'model_lr' model and the 'model_svm' along with a confusion matrix for each of them.
- ii. (1 points) Report the accuracy and classification report for each of the noisy models (model_lr_noise<nl>, model svm noise<nl>).
- iii. (1 point) Make a calibration plot for each of the models without noisy data ('model_lr', 'model_svm'). Use the CalibrationDisplay class from sklearn.calibration to make your plot.

e. Discussion (8 points)

i. (4 points) Compare the performance of the Logistic Regression (model_lr) classifier and the SVM classifier (model_svm) trained on the clean training data sets. Using the classification reports which model performs better? Mention the specific numbers (i.e., from your results calculating the precision, recall, F1 score) in a table.

| Model Name | Class | Precision | Recall | F1 |
|------------|-------|-----------|--------|------|
| model_lr | 0 | 0.90 | 0.93 | 0.91 |
| model_lr | 1 | 0.87 | 0.82 | 0.84 |
| model_svm | 0 | 0.83 | 0.98 | 0.90 |
| model_svm | 1 | 0.96 | 0.67 | 0.79 |

- ii. (4 points) Report the performance of all the classification models you have trained thus far in terms of (precision, recall, F1 scores). From the results of the three noisy models implemented using Logistic Regression.
 - 1. What did you notice about the accuracy as the noise of the dataset increased?
 - a. That the accuracy decreased
 - 2. What can you say about the effect of noise on data mining pipelines?
 - We need to minimize the impact of noise on the pipelines as it has a significant effect on the results.

| Model Name | Class | Precision | Recall | F1 |
|-------------------|--------------|-----------|--------|------|
| model_lr | Not Survived | 0.90 | 0.93 | 0.91 |
| model_lr | Survived | 0.87 | 0.82 | 0.84 |
| model_lr_noise20 | Not Survived | 0.70 | 0.94 | 0.80 |
| model lr noise20 | Survived | 0.75 | 0.32 | 0.45 |
| model_lr_noise40 | Not Survived | 0.73 | 0.77 | 0.75 |
| model_lr_noise40 | Survived | 0.57 | 0.52 | 0.54 |
| model lr noise60 | Not Survived | 0.19 | 0.03 | 0.05 |
| model_lr_noise60 | Survived | 0.33 | 0.80 | 0.46 |
| model_svm | Not Survived | 0.83 | 0.98 | 0.90 |
| model_svm | Survived | 0.96 | 0.67 | 0.79 |
| model svm noise20 | Not Survived | 0.63 | 1.00 | 0.77 |
| model_svm_noise20 | Survived | 0.00 | 0.00 | 0.00 |
| model_svm_noise40 | Not Survived | 0.63 | 1.00 | 0.77 |
| model_svm_noise40 | Survived | 0.00 | 0.00 | 0.00 |
| model_svm_noise60 | Not Survived | 0.63 | 1.0 | 0.77 |
| model_svm_noise60 | Survived | 0.00 | 0.00 | 0.00 |

Q3. Bank Churn Classification Problem (33 points)

- a. **Dataset Description:** Banking is one of those traditional industries that has gone through a steady transformation over the past few decades. Yet, many banks today with a sizeable customer base are hoping to gain a competitive edge but have not tapped into the vast amounts of data they have, especially in solving one of the most acknowledged problems *customer churn (i.e., a customer leaving the bank)*. It is advantageous to banks to know what leads a client to leave the bank. Banks often use the customer churn rate as one of their key business metrics because the cost of retaining existing customers is far less than acquiring new ones, and meanwhile increasing customer retention can greatly increase profits. *Churn prevention* allows companies to develop different programs such as loyalty and retention programs to keep as many customers as possible. Following are the attributes of the dataset we will be working with.
 - i. RowNumber (continuous) corresponds to the record (row) number and has no effect on the output.
 - ii. CustomerId (categorical)— contains random values and has no effect on customer leaving the bank.
 - iii. Surname (categorical)— the surname of a customer has no impact on their decision to leave the bank
 - iv. CreditScore (continuous) can influence customer churn, since a customer with a higher credit score is less likely to leave the bank.
 - v. Geography (categorical) a customer's location can affect their decision to leave the bank.

- vi. Gender (categorical) it's interesting to explore whether gender plays a role in a customer leaving the
- vii. Age (continuous) this is certainly relevant, since older customers are less likely to leave their bank than younger ones.
- viii. Tenure (continuous) refers to the number of years that the customer has been a client of the bank. Normally, older clients are more loyal and less likely to leave a bank.
- ix. Balance (continuous) also a very good indicator of customer churn, as people with a higher balance in their accounts are less likely to leave the bank compared to those with lower balances.
- x. NumOfProducts (continuous) refers to the number of products that a customer has purchased through the bank.
- xi. HasCrCard (categorical) denotes whether a customer has a credit card. This column is also relevant since people with a credit card are less likely to leave the bank.
- xii. IsActiveMember (categorical) active customers are less likely to leave the bank.
- xiii. EstimatedSalary (continuous) as with balance, people with lower salaries are more likely to leave the bank compared to those with higher salaries.
- xiv. Exited (Categorical) whether or not the customer left the bank. (Target variable)

b. Data Loading / Preprocessing (10 points)

- i. Loading (2 points)
 - 1. Load the data <BankChurn.csv> as a pandas dataframe using the 'pd.read_csv()' function which returns a dataframe, store this value in a variable named 'df'.
 - 2. The resulting dataframe should have the shape (10000,14) indicating that there are 10000 instances and 14 columns.
 - 3. In this dataframe, currently you have 9 features which are the following: RowNumber, CustomerID, Surname, CreditScore, Geography, Gender, Age, Tenure, Balance, NumOfProducts, HasCrCard, IsActiveMember, EstimatedSalary. Using the 'pandas.dataframe.drop' function to drop the RowNumber, CustomerID and Surname columns.
 - 4. Using the 'pandas.isnull()' function check if there are any missing values in the dataframe and report this value (i.e., the number of missing values per column of the dataframe).
 - 5. Your task is to use feature columns to predict the target column (which is categorical in our case). This can be cast as a classification problem.
 - 6. Create a dataframe X of features (by dropping the 'Exited' column from the original dataframe). Create a Pandas Series object of targets Y (by only considering the 'Exited' column from the original dataframe). Moving forward, we will be working with X and Y.
- ii. Data Visualization (4 points)
 - 1. Visualize the distribution of the 'Age' and 'CreditScore' column using the 'matplotlib.pyplot.hist' function as two separate plots. Label the x-axis and the y-axis along with giving the plot a title and assign a bin size of 7.
 - a. What are the respective mean values of these two features (use the pandas.DataFrame.mean() function)?
 - i. 38.9218 and 650.5288
 - b. What is the respective standard deviation of these two features (use the <u>pandas.DataFrame.std()</u> function)?

i. 10.49 and 96.65

- 2. **Only for this question** use the dataframe consisting of the target variable (initialized as 'df'). Using matplotlib visualize the number of males and females in each country who are active members and not active members. (Visualize this using a **barchart**. You will need to use the 'Gender', 'Geography' and 'IsActiveMember' features for this question). Visualize these graphs on two separate plots with respect to their active status. To create a barchart using matplotlib use the 'matplotlib.pyplot.bar()' function. Also label the x-axis, y-axis and give the plots a title.
 - a. How many males are from France and are active members? 1400
 - b. How many females are from Spain and are active members? 600
 - c. How many males are from France or Germany who are not active members? 1700
- 3. Using the target variable in Y plot a bar chart showing the distribution of the 'Exited' column (To create a barchart using matplotlib use the 'matplotlib.pyplot.bar()' function).
 - a. What can be said about this distribution (specifically keeping in mind this distribution represents the target variable) will this have an impact on the results of the classification model?
 - i. More people stayed than exited
 - ii. This will have an impact on the result
- 4. So far you should have successfully been able to load, preprocess and visualize your data. Now, use the 'pd.get_dummies()' function to convert categorical data into dummy variables ('Gender' and 'Geography'). (Perform this only on X).
 - a. What is the shape of X?
 - i. 10000, 11
- iii. Data Splitting (1 point)
 - 1. Split data into training and test sets using the sklearn 'train_test_split() function in a 80:20 ratio. The result of your data split should be X_train, X_test, y_train, y_test. (Respectively your training features, testing features, training targets and testing target arrays).
- iv. Data Scaling (3 points)
 - 1. Employ the 'MinMaxScaler' function on the continuous attributes in X_train. Employ the 'fit_transform()' function of the scaler to retrieve the new (scaled) version of the training data (i.e., fit transform() should be run on 'X train'). Store the result in X train again.
 - 2. Scale the X_test data using the scaler you have just *fit*, this time using the 'transform()' function. Note: store the scaled values back into X_test. At the end of this step, you must have X_train, X_test, scaled according to the MinMaxScaler.
- c. Modelling (10 points)
 - i. (2 points) Modeling (Model Instantiation / Training) using Logistic Regression classifier
 - 1. Employ the <u>Logistic Regression classifier</u> from sklearn and instantiate the model. Label/Name this model as 'model 1 lr'
 - 2. Once instantiated, `fit()` the model using the $scaled\ X_{train}$, y_{train} data.
 - 3. Employ the `predict()` function to obtain predictions on X_test and store this in a variable labeled as 'y_pred_lr'.
 - 4. Employ the 'accuracy score()' function by using the 'y_pred_lr' and 'y_test' variables as the functions parameters and print the accuracy of the Logistic Regression model
 - ii. (2 points) Modeling (Model Instantiation / Training) using Support Vector Machine Classifier

- 1. Employ the Support Vector Machine (SVM) classifier from sklearn and instantiate the model. Label/Name this model as 'model 2 svm'
- 2. Once instantiated, 'fit()' the model using the scaled X train, y train data.
- 3. Employ the 'predict()' function to obtain predictions on X_test and store this in a variable labeled as 'y pred svm'.
- Employ the <u>'accuracy_score'</u> function ('sklearn.metrics.accuracy()' function) by using the 'y_pred_lr' and 'y_test' variables as the functions parameters and print the accuracy of the SVM model.

iii. (2 points) Modeling Logistic Regression Classifier on a balanced dataset

- 1. Employ Synthetic Minority Oversampling on X_train and y_train. To use SMOTE you will have to install the imbalanced-learn library, this can either be down by executing the following command 'pip install -U imbalanced-learn' command 'conda install -c conda-forage imbalanced-learn' command for the Anaconda Cloud platform. (For more information click the following link: https://imbalanced-learn.org/stable/install.html).
 Import the 'SMOTE' function from the 'imblearn.over_sampling'. Use the 'smote.refit_resample()' function on X_train and y_train using its default parameters. Store them in X_train_smote, y_train_smote. Be careful to employ SMOTE ONLY on the training data and not on the full dataset because that can cause inadvertent "data leakage" (please see: https://arxiv.org/pdf/2107.00079.pdf for details).
- 2. Employ a new Logistic Regression classifier from *sklearn* and instantiate the model. Label this model as 'model 3 smote lr'
- 3. Once instantiated, 'fit()' the model using the balanced X train smote, y train smote data.
- 4. Employ the 'predict()' function to obtain predictions on X_test and store this in a variable labeled as 'y pred smote lr'.
- 5. Employ the 'accuracy_score' function by using the 'y_pred_lr' and 'y_test' variables as the functions parameters and print the accuracy of the new Logistic Regression model.
 - a. What is your initial observation of the accuracy of model_3_smote_lr vs. accuracy of model_1_lr? What could be the reasoning for (any possible) change in accuracy?
 - i. The accuracy of the 1_lr is higher than 3_smote_lr. This must be due to the fact that all of the models are impacted by the noise.

iv. (2 points) Modeling SVM on a balanced dataset

- Now we will employ Synthetic Minority Oversampling with an SVM classifier by training a new SVM classifier on the SMOTE balanced data.
- 2. Employ a new <u>SVM</u> classifier from *sklearn* and instantiate the model. Label this model as 'model 4 smote svm'
- 3. Once instantiated, 'fit()' the model using the *balanced* X_train_smote, y_train_smote data.
- 4. Employ the 'predict()' function to obtain predictions on X_test and store this in a variable labeled as 'y_pred_smote_svm'.
- Employ the 'accuracy_score' function ('sklearn.metrics.accuracy()' function) by using the 'y_pred_lr' and 'y_test' variables as the functions parameters and print the accuracy of the new SVM model.

- a. What is your initial observation of the accuracy of model_4_smote_svm vs. accuracy of model_2_svm? What could be the reasoning for (any possible) change in accuracy?
 - i. 0.7835 vs 0.851. Clearly smote is lower still.

v. (2 points) Modeling Grid Search Parameter Selection for SVM

- 1. We will now be reverting to our X_train and y_train data. Initialize a variable labeled as 'param grid' storing the following: {"gamma": [0.001, 0.01, 0.1], "C": [1,10,100,1000,10000]}.
- 2. Employ the <u>GridSearchCV</u> function and initialize the following parameters: estimator = SVC(), param grid = param grid, cv=5, verbose =1, scoring = 'accuracy'.
- 3. Once instantiated, 'fit()' the model using the X_train_smote, y_train_smote data. This will train SVM models with 5-fold cross validation, so it will take some time to train. Optionally, you can set the n_jobs parameter to parallelize the grid search across processes and make the fit function run faster.
- 4. Print the best paramaters using the 'best_params_' attribute and print the mean cross validated score of the best estimator (hint use the 'best score ' attribute).
- 5. Employ the 'score' function by using the 'X_test' and 'y_test' variables as the functions parameters and print the accuracy of the new gridsearch SVM model. SVM model.

d. Evaluation (5 points)

- i. (2 points) Calculate F1 Score, Precision, Recall, Accuracy (All on the test set X test, y test)
 - 1. Employ the 'classification_report()' function from sklearn.metrics to report the precision recall, fl score and accuracy for each class for the first **four** models (**parts c.i c.iv**).
- ii. (2 points) Visualize a confusion matrix for the first four models
 - Employ the `confusion_matrix()` function from sklearn.metrics to report the confusion matrix results.
 - 2. Report the False Negative and False Positive values for model 1 lr.
- iii. (1 point) Report the best F1 score of the grid search implemented in the fifth model (**part c.v**). Also report the best parameters from the grid search on the training set.

e. <u>Discussion</u> (8 points)

i. (2 points) Compare the performance of the Logistic Regression (model_1_lr) classifier and the SVM classifier (model_2_svm). Using the classification reports which model performs better? Mention the specific numbers (i.e., from your results calculating the precision, recall, F1 score) in a table.

| Model Name | Class | Precision | Recall | F1 |
|-------------|-------|-----------|--------|------|
| model_1_lr | 0 | 0.83 | 0.96 | 0.89 |
| model_1_lr | 1 | 0.57 | 0.19 | 0.29 |
| model_2_svm | 0 | 0.85 | 0.99 | 0.91 |
| model 2 svm | 1 | 0.85 | 0.30 | 0.44 |

ii. (2 points) Compare the results between the models prior to SMOTE and after. Use a combination of the classification report and the confusion matrix to emphasize your observation. (model_1_lr vs model_3_lr and model_2_svm vs model_4_svm)

| Model Name | Class | Precision | Recall | F1 |
|-------------------|-------|-----------|--------|------|
| model_1_smote_lr | 0 | 0.91 | 0.73 | 0.81 |
| model 1 smote lr | 1 | 0.39 | 0.72 | 0.51 |
| model_2_smote_svm | 0 | 0.92 | 0.78 | 0.85 |
| model 2 smote svm | 1 | 0.45 | 0.72 | 0.55 |

- iii. (2 points) Discuss the advantages and disadvantages between oversampling and under-sampling. Use the following article to help you formulate your answer and reasoning.
 (https://www.mastersindatascience.org/learning/statistics-data-science/undersampling/)
 - When a dataset is imbalanced in terms of one feature having more data than another it is helpful to balance the amount of each feature to achieve more accurate results even though there is the risk that you may lose some information from the portion of data that is lost. Two techniques for dealing with this are undersampling and oversampling. When you are lacking in information oversampling can be useful as data is generated to fill the gap. The SMOTE technique we leveraged in our notebook is one such technique for creating artificial events. When you have more than enough data for analysis undersampling is often employed. With undersampling you use all the rare events and the excess events are eliminated creating balance.
- iv. (2 points) From the results of the grid-search, comment on importance of the gamma parameter on SVM performance. What effect will the gamma value have if it is too big or too small?
 - Gamma is important because it influences each training example. When gamma is too small it
 may lead to underfitting, missing important patterns, or lead to overfitting and become too
 sensitive to noise in the training data.