Aluno: Vinícius França Lima Vilaça Regressão Logística 1. A questão 1 busca validar os dados do database utilizado. Primeiramente é necessário validar os tipos de dados, precisando que estes sejam númericos para que seja possível efetuar os cálculos corretamente. Em segundo momento se estuda a escala dos valores, onde é preciso realizar a uniformização dos dados, ou seja, colocar diferentes variáveis na mesma escala, é interessante que se faça quando estamos lidando com algoritmos baseados no Gradiente Descendente, pois estes algoritmos são sensíveis ao range dos valores dos dados e dessa forma a acurácia do algoritmo irá ser beneficiada pois o limiar de decisão da função logística ficará bem definido. Avalise também a distribuição das classes, assim pode se verificar se o dataset está desbalanceado, ou seja, se o algoritmo pode ser tendencioso ou não. Por fim, caso os dados da classicação não estejam em dados númericos é necessario que se codifique os valores para que seja possível utilizar o classificador. 2. A questão 2 realiza o feature selection, que significa buscas as features que possuem os valóres mais precisos e preciosos para o modelo. Dito isto, a correlação das features é estudada, a correlação pode ajudar a predizer um atributo de outro, ou identificar atributos que não interferem no modelo pois não possuem relação com nenhum outro. Correlação pode identificar um problema conhecido como Multicollinearity quando uma variável pode ser linearmente predita por outra com um alto grau de precisão, isso pode levar a resultados distorcidos ou enganosos e por isso e por outros problemas é um bom método de avaliação. 3. A questão 3 faz a separação dos dados de treino e teste, essa etapa é importante para evitar um modelo com underfitting ou overfitting. Foi utilizado o método Cross Validator, a validação cruzada é uma fusão de StratifiedKFold e ShuffleSplit, que retorna folds estratificados aleatórios em que os fold são feitos preservando o percentual de amostras de cada classe. 4. O modelo foi implementado ao final do notbook... 5. A acurácia foi calculada ao final do notbook... **Logistic Regression and Classification Error Metrics** Introduction We will be using the <u>Human Activity Recognition with Smartphones</u> database, which was built from the recordings of study participants performing activities of daily living (ADL) while carrying a smartphone with an embedded inertial sensors. The objective is to classify activities into one of the six activities (walking, walking upstairs, walking downstairs, sitting, standing, and laying) performed. For each record in the dataset it is provided: Triaxial acceleration from the accelerometer (total acceleration) and the estimated body acceleration. Triaxial Angular velocity from the gyroscope. A 561-feature vector with time and frequency domain variables. · Its activity label. More information about the features is available on the website: above or at https://www.kaggle.com/uciml/human-activity-recognition- with-smartphones In [48]: from __future__ import print_function #Data Path has to be set as per the file location in your system #data_path = ['..', 'data'] $data_path = ['.']$ **Question 1** Import the data and do the following: Examine the data types--there are many columns, so it might be wise to use value counts Determine if the floating point values need to be scaled Determine the breakdown of each activity Encode the activity label as an integer import pandas as pd In [50]: import numpy as np #The filepath is dependent on the data_path set in the previous cell filepath = os.sep.join(data_path + ['data.csv']) data = pd.read_csv(filepath, sep=',') Out[50]: tBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAccfBodyB max()-X mean()-Y std()-X std()-Z mad()-Y mean()-X mean()-Z std()-Y mad()-X mad()-Z -0.983111 -0.923527 0.288585 -0.020294 -0.132905 -0.995279 -0.913526 -0.995112 -0.983185 -0.934724 -0.123520 0.278419 -0.016411 -0.998245 -0.975300 -0.960322 -0.974914 -0.957686 1 -0.998807 -0.943068 0.279653 -0.978944 -0.019467 -0.113462 -0.995380 -0.967187 -0.996520 -0.963668 -0.977469 -0.938692 3 0.279174 -0.026201 -0.123283-0.996091 -0.983403 -0.997099 -0.990675 -0.982750 -0.989302 -0.938692 -0.115362 -0.998139 0.276629 -0.016570 -0.980817 -0.990482 -0.998321 -0.979672 -0.990441 -0.942469 10294 0.310155 -0.053391 -0.099109 -0.287866 -0.140589 -0.215088 -0.356083 -0.148775 -0.232057 0.185361 10295 0.363385 -0.039214 -0.105915 -0.305388 0.028148 -0.196373 -0.373540 -0.030036 -0.270237 0.185361 10296 0.349966 0.030077 -0.115788 -0.329638 -0.042143 -0.250181 -0.388017 -0.133257 -0.347029 0.007471 10297 0.237594 0.018467 -0.096499 -0.323114 -0.229775 -0.207574 -0.289477 0.007471 -0.392380 -0.279610 10298 0.153627 -0.018437 -0.137018 -0.330046 -0.195253 -0.164339 -0.430974 -0.218295 -0.229933 -0.111527 10299 rows × 562 columns The data columns are all floats except for the activity label. In [52]: data.dtypes.value_counts() Out[52]: float64 561 object dtype: int64 In [54]: data.dtypes.tail(10) Out[54]: fBodyBodyGyroJerkMag-skewness() float64 fBodyBodyGyroJerkMag-kurtosis() float64 angle(tBodyAccMean, gravity) float64 angle(tBodyAccJerkMean), gravityMean) float64 angle(tBodyGyroMean, gravityMean) float64 angle(tBodyGyroJerkMean, gravityMean) float64 angle(X, gravityMean) float64 angle(Y, gravityMean) float64 float64 angle(Z, gravityMean) object Activity dtype: object The data are all scaled from -1 (minimum) to 1.0 (maximum). In [56]: data.iloc[:, :-1].min().value_counts() Out[56]: -1.0 561 dtype: int64 In [58]: data.iloc[:, :-1].max().value_counts() Out[58]: 561 dtype: int64 Examine the breakdown of activities--they are relatively balanced. In [60]: | data.Activity.value_counts() Out[60]: LAYING 1944 **STANDING** 1906 SITTING 1777 WALKING 1722 WALKING_UPSTAIRS 1544 WALKING DOWNSTAIRS 1406 Name: Activity, dtype: int64 Scikit learn classifiers won't accept a sparse matrix for the prediction column. Thus, either LabelEncoder needs to be used to convert the activity labels to integers, or if DictVectorizer is used, the resulting matrix must be converted to a non-sparse array. Use LabelEncoder to fit transform the "Activity" column, and look at 5 random values. In [62]: from sklearn.preprocessing import LabelEncoder le = LabelEncoder() data['Activity'] = le.fit_transform(data.Activity) data['Activity'].sample(5) Out[62]: 7972 497 5 886 1 3026 5 5 4532 Name: Activity, dtype: int64 **Question 2** Calculate the correlations between the dependent variables. Create a histogram of the correlation values • Identify those that are most correlated (either positively or negatively). In [64]: # Calculate the correlation values feature_cols = data.columns[:-1] corr_values = data[feature_cols].corr() corr_values Out[64]: tBodyAcctBodyAcctBodyAcc- tBodyAcctBodyAcctBodyAcc- tBodyAcc- tBodyA mean()-X mean()-Y std()-X std()-Z mad()-X mad()-Y mean()-Z std()-Y tBodyAcc-mean()-X 1.000000 0.128037 -0.230302 0.004590 -0.016785 -0.036071 0.010303 -0.017488 -C tBodyAcc-mean()-Y 0.128037 1.000000 -0.029882 -0.046352 -0.046996 -0.054153 -0.045247 -0.047673 -C -0.022872 -0.022966 -C tBodyAcc-mean()-Z -0.230302 -0.029882 1.000000 -0.024185 -0.023745 -0.015632 0.916087 C tBodyAcc-std()-X 0.004590 -0.046352 -0.024185 1.000000 0.922525 0.861910 0.998662 tBodyAcc-std()-Y -0.016785 -0.046996 -0.023745 0.922525 1.000000 0.888259 0.918561 0.997510 C angle(tBodyGyroMean,gravityMean) 0.036047 0.013241 -0.066233 0.027464 0.001902 -0.004984 0.027729 -0.002924 -C angle(tBodyGyroJerkMean,gravityMean) 0.034296 0.077627 -0.030748 -0.027123 -0.015784 -0.012196 -0.027097 -0.013411 -C angle(X,gravityMean) -0.041021 -0.007513 0.003215 -0.374104 -0.381391 -0.353271 -0.371168 -0.378013 -C angle(Y,gravityMean) 0.034053 -0.005616 -0.012986 0.449425 0.506106 0.459092 0.444926 0.507947 C angle(Z,gravityMean) 0.030656 -0.016233 -0.028406 0.393063 0.425511 0.483424 0.389481 0.424479 561 rows × 561 columns # Simplify by emptying all the data below the diagonal tril_index = np.tril_indices_from(corr_values) tril_index Out[66]: (array([0, 1, 1, ..., 560, 560, 560]), 1, ..., 558, 559, 560])) In [68]: # Make the unused values NaNs for coord in zip(*tril_index): corr_values.iloc[coord[0], coord[1]] = np.NaN corr_values Out[68]: tBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAcctBodyAccmean()-X mean()-Y mean()-Z std()-X std()-Y std()-Z mad()-X mad()-Y -0.016785 -0.036071 0.128037 -0.230302 0.004590 0.010303 -0.017488 tBodyAcc-mean()-X NaN NaN -0.029882 -0.046352 -0.046996 -0.054153 -0.045247 -0.047673 tBodyAcc-mean()-Y NaN -C -0.024185 -0.023745 -0.015632 -0.022872 -0.022966 tBodyAcc-mean()-Z NaN NaN NaN tBodyAcc-std()-X NaN NaN NaN NaN 0.922525 0.861910 0.998662 0.916087 C NaN 0.888259 0.918561 0.997510 C tBodyAcc-std()-Y NaN NaN NaN NaN NaN angle(tBodyGyroMean,gravityMean) NaN NaN NaN NaN NaN NaN NaN angle(tBodyGyroJerkMean,gravityMean) NaN NaN NaN NaN NaN NaN NaN NaN angle(X,gravityMean) NaN angle(Y,gravityMean) NaN angle(Z,gravityMean) NaN NaN 561 rows × 561 columns # Stack the data and convert to a data frame In [70]: corr_values = (corr_values.stack().to_frame().reset_index().rename(columns={'level_0':'feature1','le vel_1':'feature2',0:'correlation'})) corr_values Out[70]: feature1 feature2 correlation tBodyAcc-mean()-X tBodyAcc-mean()-Y 0.128037 0 1 tBodyAcc-mean()-Z -0.230302 tBodyAcc-mean()-X 0.004590 2 tBodyAcc-mean()-X tBodyAcc-std()-X 3 tBodyAcc-mean()-X tBodyAcc-std()-Y -0.016785 4 tBodyAcc-mean()-X tBodyAcc-std()-Z -0.036071 -0.004582 angle(tBodyGyroJerkMean,gravityMean) angle(Y,gravityMean) angle(tBodyGyroJerkMean,gravityMean) -0.012549 angle(Z,gravityMean) 157077 -0.748249 angle(X,gravityMean) angle(Y,gravityMean) 157078 angle(X,gravityMean) angle(Z,gravityMean) -0.635231 157079 0.545614 angle(Y,gravityMean) angle(Z,gravityMean) 157080 rows × 3 columns # Get the absolute values for sorting In [71]: corr_values['abs_correlation'] = corr_values.correlation.abs() corr_values Out[71]: feature1 abs correlation feature2 correlation tBodyAcc-mean()-X 0.128037 0.128037 0 tBodyAcc-mean()-Y -0.230302 0.230302 1 tBodyAcc-mean()-Z tBodyAcc-mean()-X tBodyAcc-mean()-X tBodyAcc-std()-X 0.004590 0.004590 3 tBodyAcc-mean()-X tBodyAcc-std()-Y -0.016785 0.016785 0.036071 -0.004582 0.004582 **157075** angle(tBodyGyroJerkMean,gravityMean) angle(Y,gravityMean) 0.012549 angle(tBodyGyroJerkMean,gravityMean) angle(Z,gravityMean) -0.012549 157077 angle(X,gravityMean) angle(Y,gravityMean) -0.7482490.748249 157078 -0.635231 angle(X,gravityMean) 0.635231 angle(Z,gravityMean) 157079 angle(Z,gravityMean) 0.545614 angle(Y,gravityMean) 0.545614 157080 rows × 4 columns A histogram of the absolute value correlations. In [72]: import matplotlib.pyplot as plt import seaborn as sns %matplotlib inline In [73]: sns.set_context('talk') sns.set_style('white') sns.set_palette('dark') ax = corr_values.abs_correlation.hist(bins=50) ax.set(xlabel='Absolute Correlation', ylabel='Frequency') Out[73]: [Text(0.5, 0, 'Absolute Correlation'), Text(0, 0.5, 'Frequency')] 10000 8000 Frequency 6000 4000 2000 0 0.0 0.2 0.4 0.6 0.8 1.0 **Absolute Correlation** In [74]: # The most highly correlated values corr_values.sort_values('correlation', ascending=False).query('abs_correlation>0.8') Out[74]: feature1 feature2 correlation abs_correlation 156894 1.000000 fBodyBodyGyroJerkMag-mean() fBodyBodyGyroJerkMag-sma() 1.000000 93902 tBodyAccMag-sma() tGravityAccMag-sma() 1.000000 1.000000 101139 1.000000 1.000000 tBodyAccJerkMag-mean() tBodyAccJerkMag-sma() 96706 tGravityAccMag-mean() tGravityAccMag-sma() 1.000000 1.000000 94257 tBodyAccMag-energy() tGravityAccMag-energy() 1.000000 1.000000 0.993425 22657 tGravityAcc-mean()-Y angle(Y,gravityMean) -0.993425 39225 tGravityAcc-arCoeff()-Z,3 tGravityAcc-arCoeff()-Z,4 38739 tGravityAcc-arCoeff()-Z,2 tGravityAcc-arCoeff()-Z,3 -0.994628 0.994628 23176 tGravityAcc-mean()-Z angle(Z,gravityMean) -0.994764 0.994764 38252 tGravityAcc-arCoeff()-Z,1 tGravityAcc-arCoeff()-Z,2 -0.995195 0.995195 22815 rows × 4 columns **Question 3** Split the data into train and test data sets. This can be done using any method, but consider using Scikit-learn's StratifiedShuffleSplit to maintain the same ratio of predictor classes. Regardless of methods used to split the data, compare the ratio of classes in both the train and test splits. from sklearn.model_selection import StratifiedShuffleSplit # Get the split indexes strat_shuf_split = StratifiedShuffleSplit(n_splits=1, test_size=0.3, random_state=42) train_idx, test_idx = next(strat_shuf_split.split(data[feature_cols], data.Activity)) # Create the dataframes X_train = data.loc[train_idx, feature_cols] y_train = data.loc[train_idx, 'Activity'] X_test = data.loc[test_idx, feature_cols] y_test = data.loc[test_idx, 'Activity'] In [76]: y_train.value_counts(normalize=True) Out[76]: 0 0.188792 0.185046 1 0.172562 3 0.167152 5 0.149951 0.136496 Name: Activity, dtype: float64 In [77]: y_test.value_counts(normalize=True) Out[77]: 0 0.188673 0.185113 0.172492 0.167314 5 0.149838 0.136570 Name: Activity, dtype: float64 **Question 4** • Fit a logistic regression model without any regularization using all of the features. Be sure to read the documentation about fitting a multi-class model so you understand the coefficient output. Store the model. In [78]: **from sklearn.linear_model import** LogisticRegression # Standard logistic regressio model_lr = LogisticRegression().fit(X_train, y_train) In [79]: predicted_activity = model_lr.predict(X_test)

Question 5

accuracy

Out[80]: 0.9805825242718447

In [80]:

In []:

Calculate the following error metric:

accuracy = sum(predicted_activity==y_test) / len(y_test)