Human Activity Recognition Using Smartphones Data Set

Introduction

We will be using the Human Activity Recognition with Smartphones 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.

Plan for data exploration

- 1. Exploring data
 - Examine the data types and value_counts
- 2. feature engineering
 - see the data distribution
 - removing unimportant data if found
 - · dealing with missing (NaN) values if found
 - feature scalling for continuous variables if needed
- 3. encoding
 - encoding for categorical variables if found as to Encode the activity label as an integer
- 4. Spliting the Data
 - StratifiedShuffleSplit
- 5. Applying classification models
 - Logistic Regression
 - K-Nearest Neighbors (KNeighbors)
 - Decision Trees
 - Ensemble Methods (Gradient Boosting)
- 6. Selecting the best model
- 7. Next steps

Exploring and feature engineering

```
# importing
          import pandas as pd
          import numpy as np
          import matplotlib.pyplot as plt
          import seaborn as sns
          from sklearn.model_selection import KFold, cross_val_predict, GridSearchCV, train_te
          from sklearn.metrics import r2_score, mean_squared_error, mean_absolute_error
          from sklearn.pipeline import Pipeline
          %matplotlib inline
In [ ]:
          filepath = 'data/Human_Activity_Recognition_Using_Smartphones_Data.csv'
          data = pd.read_csv(filepath, sep=',')
          data.head()
            tBodyAcc-
                      tBodyAcc-
                                 tBodyAcc-
                                            tBodyAcc- tBodyAcc-
                                                                  tBodyAcc- tBodyAcc-
                                                                                       tBodyAcc-
Out[]:
                                                          std()-Y
             mean()-X
                        mean()-Y
                                   mean()-Z
                                               std()-X
                                                                     std()-Z
                                                                               mad()-X
                                                                                         mad()-Y
                                                                                                    n
         0
             0.288585
                        -0.020294
                                  -0.132905
                                             -0.995279
                                                        -0.983111
                                                                   -0.913526
                                                                              -0.995112
                                                                                         -0.983185
                                                                                                    -0
         1
             0.278419
                        -0.016411
                                  -0.123520
                                                        -0.975300
                                                                   -0.960322
                                                                                                    -0
                                             -0.998245
                                                                              -0.998807
                                                                                         -0.974914
         2
              0.279653
                        -0.019467
                                  -0.113462
                                             -0.995380
                                                        -0.967187
                                                                   -0.978944
                                                                              -0.996520
                                                                                         -0.963668
                                                                                                    -0
             0.279174
         3
                        -0.026201
                                  -0.123283
                                             -0.996091
                                                        -0.983403
                                                                   -0.990675
                                                                              -0.997099
                                                                                         -0.982750
                                                                                                   -0
         4
              0.276629
                        -0.016570
                                  -0.115362
                                             -0.998139
                                                        -0.980817
                                                                   -0.990482
                                                                              -0.998321
                                                                                         -0.979672
                                                                                                    -0
        5 rows × 562 columns
In [ ]:
          data.dtypes.value_counts()
         float64
                     561
Out[]:
         object
                       1
         dtype: int64
        The data columns are all floats except for the activity label.
In [ ]:
          # see the min and max of data excluding the target
          print('min = ',data.iloc[:, :-1].min().value_counts())
          print('min = ',data.iloc[:, :-1].max().value_counts())
         min = -1.0
                         561
         dtype: int64
         min = 1.0
                        561
         dtype: int64
        The data are all scaled from -1 (minimum) to 1.0 (maximum).
In [ ]:
          # Examine the breakdown of activities -- to see if balanced or not
          data.Activity.value_counts(normalize=True)
         LAYING
                                 0.188756
Out[]:
                                 0.185067
         STANDING
         SITTING
                                 0.172541
         WALKING
                                 0.167201
         WALKING UPSTAIRS
                                 0.149917
         WALKING_DOWNSTAIRS
                                 0.136518
         Name: Activity, dtype: float64
```

```
In [ ]: data.isnull().sum().all()
Out[ ]: False
```

encoding

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.

we will use LabelEncoder to fit_transform the "Activity" column, and look at 5 random values.

```
In [ ]:
        from sklearn.preprocessing import LabelEncoder
         le = LabelEncoder()
         data['Activity'] = le.fit_transform(data.Activity)
         data['Activity'].sample(5)
         ### END SOLUTION
        2487
                3
Out[]:
        7854
                1
        5771
             0
        2393
               2
        1222
        Name: Activity, dtype: int32
```

looking at correlation

- 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 [ ]:
         # Calculate the correlation values
         feature cols = data.columns[:-1]
         corr_values = data[feature_cols].corr()
         # Simplify by emptying all the data below the diagonal
         tril_index = np.tril_indices_from(corr_values)
         # Make the unused values NaNs
         for coord in zip(*tril_index):
             corr values.iloc[coord[0], coord[1]] = np.NaN
         # Stack the data and convert to a data frame
         corr_values = (corr_values
                        .stack()
                        .to_frame()
                        .reset_index()
                         .rename(columns={'level_0':'feature1',
                                          'level 1': 'feature2',
                                          0:'correlation'}))
         # Get the absolute values for sorting
         corr_values['abs_correlation'] = corr_values.correlation.abs()
         corr_values
```

Out[]:		feature1	feature2	correlation	abs correlation
	0	tBodyAcc-mean()-X	tBodyAcc-mean()-Y	0.128037	0.128037
	1	tBodyAcc-mean()-X	tBodyAcc-mean()-Z	-0.230302	0.230302
	2	tBodyAcc-mean()-X	tBodyAcc-std()-X	0.004590	0.004590
	3	tBodyAcc-mean()-X	tBodyAcc-std()-Y	-0.016785	0.016785
	4	tBodyAcc-mean()-X	tBodyAcc-std()-Z	-0.036071	0.036071
		•	,		
1570 ⁻	 75	angle(tBodyGyroJerkMean,gravityMean)	angle(Y,gravityMean)	-0.004582	0.004582
1570		angle(tBodyGyroJerkMean,gravityMean)	angle(Z,gravityMean)	-0.012549	0.012549
1570		angle(X,gravityMean)	angle(Y,gravityMean)	-0.748249	0.748249
1570		angle(X,gravityMean)	angle(Z,gravityMean)	-0.635231	0.635231
1570	10	angle(x,gravityiviean)	angle(z,gravitylviean)	-0.033231	0.053251

157080 rows × 4 columns

157079

angle(Y,gravityMean) angle(Z,gravityMean)

0.545614

0.545614

Out[]:		feature1	feature2	correlation	abs_correlation
	156894	fBodyBodyGyroJerkMag-mean()	fBodyBodyGyroJerkMag-sma()	1.000000	1.000000
	93902	tBodyAccMag-sma()	tGravityAccMag-sma()	1.000000	1.000000
	101139	tBodyAccJerkMag-mean()	tBodyAccJerkMag-sma()	1.000000	1.000000
	96706	tGravityAccMag-mean()	tGravityAccMag-sma()	1.000000	1.000000
	94257	tBodyAccMag-energy()	tGravityAccMag-energy()	1.000000	1.000000
	•••				
	22657	tGravityAcc-mean()-Y	angle(Y,gravityMean)	-0.993425	0.993425
	39225	tGravityAcc-arCoeff()-Z,3	tGravityAcc-arCoeff()-Z,4	-0.994267	0.994267
	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

Data split

• This can be done using any method, but consider using Scikit-learn's StratifiedShuffleSplit to maintain the same ratio of predictor classes.

```
from sklearn.model_selection import StratifiedShuffleSplit
X = data[feature_cols]
y = data['Activity']
```

```
strat_shuf_split = StratifiedShuffleSplit(n_splits=1,
                                                    test_size=0.3,
                                                    random_state=42)
         train_idx, test_idx = next(strat_shuf_split.split(X, y))
         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 [ ]:
         y_train.value_counts(normalize=True)
             0.188792
Out[]:
        2
             0.185046
        1
             0.172562
         3
            0.167152
            0.149951
        4 0.136496
        Name: Activity, dtype: float64
In [ ]:
         y_test.value_counts(normalize=True)
Out[ ]: 0
             0.188673
        2
             0.185113
             0.172492
        1
        3
           0.167314
            0.149838
             0.136570
        Name: Activity, dtype: float64
        we maintained the distribution of the target class seccussfuly
```

Applying classification models

- Logistic Regression
- K-Nearest Neighbors (KNeighbors)
- Decision Trees
- Ensemble Methods (Gradient Boosting)

1.Logistic Regression

```
In [ ]: from sklearn.linear_model import LogisticRegression
    # Standard Logistic regression
    lr = LogisticRegression(solver='liblinear').fit(X_train, y_train)

In [ ]: from sklearn.linear_model import LogisticRegressionCV
    # L1 regularized Logistic regression
    lr_l1 = LogisticRegressionCV(Cs=10, cv=4, penalty='l1', solver='liblinear').fit(X_tr
    # L2 regularized Logistic regression
    lr_l2 = LogisticRegressionCV(Cs=10, cv=4, penalty='l2', solver='liblinear').fit(X_tr
```

- Predict and store the class for each model.
- Store the probability for the predicted class for each model.

```
In [ ]: # Predict the class and the probability for each
    y_pred = list()
    y_prob = list()

coeff_labels = ['lr', 'l1', 'l2']
    coeff_models = [lr, lr_l1, lr_l2]

for lab,mod in zip(coeff_labels, coeff_models):
        y_pred.append(pd.Series(mod.predict(X_test), name=lab))
        y_prob.append(pd.Series(mod.predict_proba(X_test).max(axis=1), name=lab))

y_pred = pd.concat(y_pred, axis=1)
    y_prob = pd.concat(y_prob, axis=1)
    y_pred.head()
```

```
In [ ]: y_prob.head()
```

```
        Out[]:
        Ir
        I1
        I2

        0
        0.998939
        0.998999
        0.999998

        1
        0.988165
        0.999272
        0.999477

        2
        0.987592
        0.995039
        0.999697

        3
        0.981381
        0.999174
        0.999865

        4
        0.998277
        0.999920
        0.999997
```

error metrics

For each model, calculate the following error metrics:

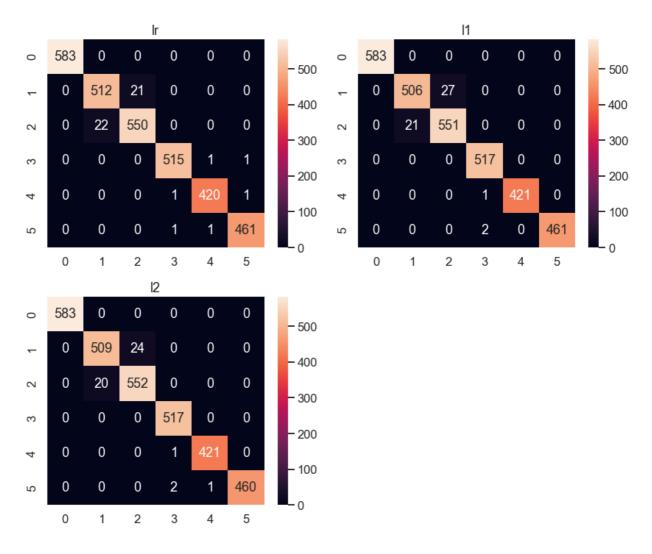
- Accuracy
- Precision
- Recall
- F-score
- Confusion Matrix

```
In []:
    from sklearn.metrics import precision_recall_fscore_support as score
    from sklearn.metrics import confusion_matrix, accuracy_score, roc_auc_score
    from sklearn.preprocessing import label_binarize

    metrics = list()
    cm = dict()

    for lab in coeff_labels:
```

```
precision, recall, fscore, _ = score(y_test, y_pred[lab], average='weighted')
              accuracy = accuracy_score(y_test, y_pred[lab])
              # ROC-AUC scores can be calculated by binarizing the data
              auc = roc_auc_score(label_binarize(y_test, classes=[0,1,2,3,4,5]),
                        label_binarize(y_pred[lab], classes=[0,1,2,3,4,5]),
                        average='weighted')
              cm[lab] = confusion_matrix(y_test, y_pred[lab])
             metrics.append(pd.Series({'precision':precision, 'recall':recall,
                                         'fscore':fscore, 'accuracy':accuracy,
                                         'auc':auc},
                                       name=lab))
         metrics = pd.concat(metrics, axis=1)
In [ ]:
         metrics
                                11
                                         12
Out[]:
                       Ir
         precision 0.984144 0.983514 0.984477
            recall 0.984142 0.983495 0.984466
           fscore 0.984143 0.983492 0.984464
         accuracy 0.984142 0.983495 0.984466
             auc 0.990384 0.989949 0.990553
In [ ]:
         f1_lr = f1_score(y_pred, y_test, average='weighted')
         f1_lr
        0.9679843882037947
Out[ ]:
In [ ]:
         ### Display or plot the confusion matrix for each model.
         fig, axList = plt.subplots(nrows=2, ncols=2)
         axList = axList.flatten()
         fig.set_size_inches(12, 10)
         axList[-1].axis('off')
         for ax,lab in zip(axList[:-1], coeff_labels):
              sns.heatmap(cm[lab], ax=ax, annot=True, fmt='d');
              ax.set(title=lab);
         plt.tight layout()
```



2.K-Nearest Neighbors (KNeighbors)

```
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import f1_score

knn = KNeighborsClassifier(n_neighbors=3, weights='distance')
knn = knn.fit(X_train, y_train)
y_pred = knn.predict(X_test)
f1_knn = f1_score(y_pred, y_test, average='weighted')
f1_knn
```

Out[]: 0.9679843882037947

3.Decision Trees

```
print('GR.best_estimator_: ', GR.best_estimator_)
         print('GR.best_score_: ', GR.best_score_)
         print('GR.best_params_: ', GR.best_params_)
        GR.best_estimator_: DecisionTreeClassifier(max_depth=2, min_samples_leaf=2, random_
        state=42)
        GR.best_score_: nan
        GR.best_params_: {'max_depth': 2, 'min_samples_leaf': 2, 'min_samples_split': 2}
In [ ]:
        GR.best_score_, GR.best_params_
        (nan, {'max_depth': 2, 'min_samples_leaf': 2, 'min_samples_split': 2})
Out[ ]:
In [ ]:
         y_pred = GR.predict(X_test)
         f1_dt = f1_score(y_pred, y_test, average='weighted')
         f1_dt
        0.6770962196812663
Out[ ]:
```

4.GradientBoosting

```
In [ ]:
         from sklearn.ensemble import GradientBoostingClassifier
         GBC = GradientBoostingClassifier(max_features=5, n_estimators=100, random_state=42)
         GBC.fit(X_train.values, y_train.values)
         y_pred = GBC.predict(X_test)
         f1_GBC = f1_score(y_pred, y_test, average='weighted')
         f1_GBC
```

0.9805798565748332 Out[]:

Selecting best model

```
In [ ]:
         pd.DataFrame({'Logistic Regression':f1_lr, 'KNN':f1_knn,
                                          'Decision Trees':f1_dt, 'Gradient Boosting':f1_GBC},in
Out[]:
                   Logistic Regression
                                        KNN Decision Trees Gradient Boosting
         F1_SCORE
                            0.967984 0.967984
                                                   0.677096
                                                                     0.98058
```

so we will chooce GradientBoostingClassifier

key findings

GradientBoostingClassifier got us the highest F1_Score so we will choose it and Decision Trees took too long and got the worst score

Next steps

we can use another encoding method like LabelBinarizer and we can try another ensemble method like Random forest