FE 520 - Into to Python for Financial Applications

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Q1. Time Series Data Practice

We start by importing numpy and pandas as two standard packages to solve this problem. Next, we import *Energy.xlsx* as pandas data-frame. We then define a function **splitEnergy()** which takes in two values *StartYear* and *EndYear*. Both have default values *2012* and *None* respectively. We start the function definition by setting the 'Data Date' column of the data-frame to type string for easy manipulation and then splitting just the year and converting it back to integer again. The function definition has one if-else as follows -

- If the EndYear is None then the TestData gets all the values with StartYear and TrainData gets the rest
 - By default, all company Data Date within 2012 will be selected as Testing data
- Else TestData gets all values within range of StartYear and EndYear and TrainData gets everything else.

TestData and TrainData are then only given columns from Accumulated Other Comprehensive Income (Loss) to Selling, General and Administrative Expenses by getting their column index by name and then dropping every other column which is not in that range.

The function then returns TestData.values and TrainData.values which in the form of numpy array.

Results

```
1 splitEnergy(2012,2013)
(array([[-1057., 1421., 8212., ...,
                                           0.,
                                                   0.,
                                                          nan],
       [ -779., 1699., 7925., ...,
                                           0.,
                                                   0.,
                                                          nan],
        [ -675., 2167., 8157., ...,
                                                   0.,
                                                          nan],
                                           0.,
        [ -254.,
                   190., 14252., ...,
                                           0.,
                                                  nan,
                                                          nanl,
        [-205.,
                   206., 13531., ...,
                                           0.,
                                                  nan,
                                                           nan],
                   197., 12737., ...,
       [-204.,
                                           0.,
                                                  nan,
                                                          nan]]),
array([[-1749.,
                  1502., 8780., ...,
                                           0.,
                                                   0.,
                                                          nan],
                          8296., ...,
        [-1603., 1434.,
                                           0.,
                                                   0.,
                                                          nan],
        [-1377.,
                   865.,
                          8839., ...,
                                           0.,
                                                   0.,
                                                          nan],
        [ -321.,
                   142., 10304., ...,
                                          nan,
                                                          nan],
                   176., 9545., ...,
        [ -327.,
                                          nan,
                                                  nan,
                                                          nan],
        [-234.,
                   236., 10401., ...,
                                          nan,
                                                  nan,
                                                          nan]]))
```

Final output of Q1

Q3. Regression

3.1 In this question, we are going to use the diabetes data set. Use sklearn.datasets.load diabetes() to load the data and labels.

We import the diabetes datasets in two variables X and y by passing the argument $return_X_y=True$ in **sklearn.datasets.load_diabetes()**

3.2. Randomly split the data into training set (80%) and testing set (20%)

Here we use train_test_split from sklearn. We pass two specific arguments in the function -

- test_size = 0.2 which gives 20% to test data and 80% to training data
- random_state = 42 which controls the shuffling(to randomize) applied to the data before applying the split.

3.3. Create a linear regression model using sklearn, and fit training data. Evaluate your model using test data. Give all the coefficient and R-squared score.

We import Linear Regression from sklearn.model. We then pass the training data (diabetes_X_train, diabetes_y_train) for the algorithm to train on (through the fit function) and then perform a prediction (using the predict function) on the testing data (diabetes_X_test). We then use the .coef_ function of the algorithm to fetch all the coefficients and finally compare the prediction with our sample space (diabetes_y_test) to calculate the R-squared score.

Result

```
Coefficients:
[ 37.90031426 -241.96624835 542.42575342 347.70830529 -931.46126093 518.04405547 163.40353476 275.31003837 736.18909839 48.67112488]
Coefficient of determination: 0.45
```

3.4. Use 10-fold cross validation to fit and validate your linear regression models on the whole data set. Print the scores for each validation.

We run K-Cross Validation (k=10) on out Linear Regression model and print the CV score for each iteration. The final output is the mean of the absolute scores (accuracy) along with the standard deviation.

Result

```
Cross Validation Score for fold 1 is: 0.5310818612097871
Cross Validation Score for fold 2 is: 0.48929200594262445
Cross Validation Score for fold 3 is: 0.6155148865078137
Cross Validation Score for fold 4 is: -0.07802446434722854
Cross Validation Score for fold 5 is: 0.4650954815453411
Cross Validation Score for fold 6 is: 0.5338119174559128
Cross Validation Score for fold 7 is: 0.706563282930452
Cross Validation Score for fold 8 is: 0.5253790639706339
Cross Validation Score for fold 9 is: -0.23181569661350765
Cross Validation Score for fold 10 is: 0.37529064060113176

Mean Accuracy: 0.46 with STD DEVIATION: 0.58
```

3.5. Use sklearn to create RandomForestRegressor model, fit the training data into it.

We import RandomForestRegressor from sklearn and pass it the following arguments -

- n estimators=1000 The number of trees in the forest.
- max_depth=None The maximum depth of the tree. If None, then nodes are expanded until all leaves are pure or until all leaves contain less than min samples split samples.
- min_samples_split=2 The minimum number of samples required to split an internal node
- random_state=1 Controls both the randomness of the bootstrapping of the samples used when building trees

We then calculate and print the R-squared score and Root mean squared test error

Result

```
R-squared score: 0.44
Root mean squared test error = 54.465358204535065
```

2.6. Use Grid Search to find the optimal hyper-parameters (max depth:{None, 7, 4} and min samples split: {2, 10, 20}) for RandomForestRegressor.

We start by setting up the **RandomForestRegressor** with the values used in the previous question. We then setup **GridSearchCV** with the parameters mentioned in the question and cv=10 and train the model.

The model gives best max_depth = None and min_samples_split = 20

Result

```
{'bootstrap': True,
 'ccp alpha': 0.0,
 'criterion': 'mse',
'max_depth': None,
'max features': 'auto',
'max leaf nodes': None,
'max samples': None,
'min impurity decrease': 0.0,
 'min impurity split': None,
 'min samples leaf': 1,
 'min samples split': 20,
 'min weight fraction leaf': 0.0,
'n estimators': 1000,
 'n jobs': None,
 'oob score': False,
 'random state': 1,
 'verbose': 0,
 'warm start': False}
```

We further calculate the *R*-squared score using the optimal parameters which comes out to be 0.46 which is better than RandomForestRegressor without GridSearch (0.44)

CODE APPENDIX

Q1

```
#!/usr/bin/env python
    # coding: utf-8
    # In[223]:
    #import statements
    import pandas as pd
    import numpy as np
    # In[224]:
    #import Energy.xlsx to pandas dataframe
    df = pd.read excel (r'Energy.xlsx')
    print(df.shape) #check if data is imported correctly by checking dimensions
    # In[225]:
    #getting column index by column name
    index 1 = df.columns.get loc("Accumulated Other Comprehensive Income (Loss)")
    index_2 = df.columns.get_loc("Selling, General and Administrative Expenses")
    # In[226]:
    #function definition for splitEnergy
    def splitEnergy(StartYear=2012,EndYear=None):
        df temp = df
        #set column 'Data Date' type to string to split year and convert back to
integer
        df temp['Data Date'] = df temp['Data Date'].astype(str).str[:4].astype(int)
        #If EndYear is None, we will only choose all data with "Data Date" ==
StartYear as Test data, all other data as Train data.
```

```
#By default, all company Data Date within 2012 will be selected as Testing
data.
        if (EndYear == None):
            TestData = df temp[df temp['Data Date'] == StartYear]
            TrainData = df temp[df temp['Data Date'] != StartYear]
        #If EndYear is NOT None, we will choose all data with "Data Date" == StartYear
to EndYear as Test data, all other data as Train data
            TestData = df temp[(df temp['Data Date'] >= StartYear) & (df temp['Data
Date'] <= EndYear)]</pre>
            TrainData = df temp[(df temp['Data Date'] < StartYear) | (df temp['Data</pre>
Date'] > EndYear)]
        #array from column "Accumulated Other Comprehensive Income (Loss)" to column "
Selling, General and Administrative Expenses".
        TestData = TestData.iloc[:, index 1:index 2]
        TrainData = TrainData.iloc[:, index_1:index_2]
        #Output Data type: Array(Numpy)
        return TestData.values , TrainData.values
    # In[227]:
    #function execution
    splitEnergy(2012,2013)
```

```
#!/usr/bin/env python
    # coding: utf-8
    # ### import statements
    # In[18]:
    import matplotlib.pyplot as plt
    import numpy as np
    from sklearn import datasets, linear model
    from sklearn.metrics import r2 score
    from sklearn.model selection import cross val score
    from sklearn.ensemble import RandomForestRegressor
    from sklearn.model selection import train test split
    from sklearn.metrics import mean squared error
    from sklearn.model_selection import GridSearchCV
    from pprint import pprint
    # ### Question 3 Regression
    # #### 3.1 & 3.2
    # In[19]:
    # Load the diabetes dataset
    diabetes_X, diabetes_y = datasets.load_diabetes(return_X_y=True)
    # Split the data into training/testing sets
    # Split the targets into training/testing sets
    diabetes X train, diabetes X test, diabetes y train, diabetes y test =
train_test_split(diabetes_X, diabetes_y, test_size=0.20, random_state=42)
    # #### 3.3
    # In[20]:
```

```
# Create linear regression object
    linreg = linear model.LinearRegression()
    # Train the model using the training sets
    linreg.fit(diabetes X train, diabetes y train)
    # Make predictions using the testing set
    diabetes y pred = linreg.predict(diabetes X test)
    # In[21]:
    # The coefficients
    print('Coefficients: \n', linreg.coef)
    # The coefficient of determination: 1 is perfect prediction
    print('\nR-squared score: %.2f'
          % r2 score(diabetes y test, diabetes y pred))
    # #### 3.4
    # In[22]:
    #Using 10-fold cross validation to fit and validate linear regression models on
the whole data set. Printing the scores for each validation.
    scores = cross val score(linreg, diabetes X train, diabetes y train, cv=10)
    count = 1
    for i in scores:
        print("Cross Validation Score for fold ", count, "is: ", i)
        count = count + 1
    print("\nMean Accuracy: %0.2f with STD DEVIATION: %0.2f" %
(np.mean(np.abs(scores)), scores.std() * 2))
    # #### 3.5
```

```
# In[26]:
    # Use sklearn to create RandomForestRegressor model, and fit the training data
into it.
    rforest = RandomForestRegressor(n estimators=1000, max depth=None,
min samples split=2, random state=1)
    rforest.fit(diabetes X train, diabetes y train)
    diabetes y pred = rforest.predict(diabetes X test)
    #calculating the r square score
    print('\nR-squared score: %.2f'% r2 score(diabetes y test, diabetes y pred))
    #calculating root mean sq error
    print("Root mean squared test error =
{0}".format(np.sqrt(np.mean((rforest.predict(diabetes X test) -
diabetes y test)**2))))
    # #### 3.6
    # In[24]:
    #Using Grid Search to find the optimal hyper-parameters
    #setup randomforest
    rforest = RandomForestRegressor(n estimators=1000, max depth=None,
min samples split=2, random state=1)
    #setup and train GridSearchCV
    clf = GridSearchCV(rforest, {'max depth': [None, 7, 4],'min samples split': [2,
10, 20]}, cv=10)
    model = clf.fit(diabetes X train, diabetes y train)
    # print optimal hyper-parameters
    pprint(model.best_estimator_.get_params())
    # In[25]:
```

```
#print R-squared score
diabetes_y_pred = model.predict(diabetes_X_test)
print('\nR-squared score: %.2f'% r2_score(diabetes_y_test, diabetes_y_pred))
```