Documentation Part2 - Πρόβλεψη, Ταξινόμηση, Ομαδοποίηση σε Καρδιοτοκογραφικά Δεδομένα

Σημείωση:

- Ο κώδικας που θα ακολουθήσει βρίσκεται στο φάκελο:
 - Code/Part2.ipynb
- Τα δεδομένα(data) βρίσκονται στο φάκελο:
 - Data/data2.csv
- Τα δεδομένα έχουν τροποποιηθεί στο αρχείο csv. Αρχικά έχουν διαγραφεί οι επικεφαλίδες έτσι ώστε να έχουμε το πρόβλημα supervised.

Επεξήγηση Κώδικα

Ερώτημα 1 (Data Cleaning):

3

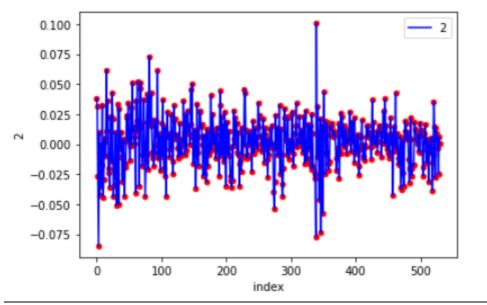
```
data = pd.read_csv("../Data/data 2.csv", header = None, sep=',')
: # Data Cleaning
  data = data[data.columns[1:]]
  data.dropna(inplace=True)
  data.reset_index(inplace=True)
  data = data.iloc[:530]
  data.info()
  print("\n")
  plt.plot(data[1])
  plt.show()
  print("\n")
  data.head()
  <class 'pandas.core.frame.DataFrame'>
  RangeIndex: 530 entries, 0 to 529
  Data columns (total 10 columns):
  # Column Non-Null Count Dtype
     index 530 non-null int64
             530 non-null float64
             530 non-null
   2
                           float64
                          float64
             530 non-null
   3
      3
   4
      4
            530 non-null float64
             530 non-null
             530 non-null float64
             530 non-null
                           float64
   8
      8
             530 non-null
                           float64
             530 non-null
                           float64
  \texttt{dtypes: float64(9), int64(1)}
  memory usage: 41.5 KB
   0.06
   0.04
   0.02
   0.00
  -0.02
  -0.04
  -0.06
               100
                      200
                              300
                                     400
                                            500
    index
                1
                       2
                                3
                                        4
                                                5
                                                         6
         0
```

1 0.025426 0.031813 0.007787 0.008455 0.012866 0.004162 0.018920 0.011341 0.008773 2 -0.028862 -0.026353 -0.030469 -0.017833 -0.028735 0.017293 -0.035899 -0.017073 -0.020015

3 -0.062208 -0.084716 0.003391 -0.011726 -0.000466 -0.040061 0.028283 -0.005561 -0.019424 4 0.009860 0.009658 -0.021533 -0.019873 -0.012710 -0.004474 -0.009764 -0.010989 -0.007802

Δημιουργία Scatter Plot για την καλύτερη αντίληψη του γραφήματος:

```
# a scatter plot comparing num_children and num_pets
ax = plt.gca()
data.plot(kind='line', x='index',y=2, color='blue', ax=ax)
data.plot(kind='scatter', x='index',y=2, color='red', ax=ax)
plt.show()
```



Καθορίζουμε το input ως προετοιμασία για την υλοποίηση τουπροβλήματος supervised:

```
#Convert data to input-output
df = data[1].to_frame(name='ISE_TL')
df.head()
```

ISE_TL

- 0.035754
- 1 0.025426
- 2 -0.028862
- 3 -0.062208
- 4 0.009860

Κανονικοποίηση Δεδομένων:

Εκτελούμε τη συνάρτηση 'timeseries_to_supervised', η οποία λύνει το πρόβλημα του supervised:

```
n_in = 6
n_out = 1
sdf = timeseries_to_supervised(df, n_in, n_out)
sdf.head()
```

	ISE_TL(t-6)	ISE_TL(t-5)	ISE_TL(t-4)	ISE_TL(t-3)	ISE_TL(t-2)	ISE_TL(t-1)	ISE_TL(t)
6	0.746889	0.668147	0.254242	0.000000	0.549467	0.251732	0.592052
7	0.668147	0.254242	0.000000	0.549467	0.251732	0.592052	0.160419
8	0.254242	0.000000	0.549467	0.251732	0.592052	0.160419	0.479339
9	0.000000	0.549467	0.251732	0.592052	0.160419	0.479339	0.642311
10	0.549467	0.251732	0.592052	0.160419	0.479339	0.642311	0.301278

timeseries to supervised:

```
def timeseries_to_supervised(df, n_in, n_out):
    agg = pd.DataFrame()

for i in range(n_in, 0, -1):
    df_shifted = df.shift(i).copy()
    df_shifted.rename(columns=lambda x: ('%s(t-%d)' % (x, i)), inplace=True)
    agg = pd.concat([agg, df_shifted], axis=1)

for i in range(0, n_out):
    df_shifted = df.shift(-i).copy()
    if i == 0:
        df_shifted.rename(columns=lambda x: ('%s(t)' % (x)), inplace=True)
    else:
        df_shifted.rename(columns=lambda x: ('%s(t+%d)' % (x, i)), inplace=True)
        agg = pd.concat([agg, df_shifted], axis=1)
    agg.dropna(inplace=True)
    return agg
```

Προετοιμασία για την εκπαίδευση:

```
X, y = sdf[('ISE_TL(t-%d)' % i)  for i in range(6, 0, -1)]].values, sdf['ISE_TL(t)'].values
print(X.shape, y.shape)
(524, 6) (524,)
# Preparation for Training
# Split data into train and test
len_data = X.shape[0]
train_size = int(len_data * .5)
print ("Train size: %d" % train_size)
print ("Test size: %d" % (len_data - train_size))
xtr, ytr = X[:train_size, :], y[:train_size]
xte, yte = X[train_size:, :], y[train_size:]
print(xtr.shape, ytr.shape)
print(xte.shape, yte.shape)
Train size: 262
Test size: 262
(262, 6) (262,)
(262, 6) (262,)
```

Ερώτημα 2α (υλοποίηση MPL):

```
----- MLP & RNN -----
batch_size = 1 #1
model = Sequential()
# MLP
xtr, ytr = X[:train_size, :], y[:train_size]
xte, yte = X[train_size:, :], y[train_size:]
model.add(Dense(units=100, input_dim=xtr.shape[1], activation="relu"))
model.add(Dense(50, activation="relu"))
model.add(Dense(1))
model.compile(loss='mean_squared_error', optimizer='adam')
model.summary()
# # RNN
# model.add(SimpleRNN(units=100, batch_input_shape=(batch_size, xtr.shape[1],
                                               xtr.shape[2]), activation="relu",
                                               return sequences=True))
# model.add(Dense(50, activation="relu"))
# model.add(Dense(1))
# model.compile(loss='mean_squared_error', optimizer='adam')
# model.summary()
model.fit(xtr,ytr, epochs=10, batch_size=batch_size, verbose=0)
4
```

Model: "sequential"

Layer (type)	Output Shape	Param #
dense (Dense)	(None, 100)	700
dense_1 (Dense)	(None, 50)	5050
dense_2 (Dense)	(None, 1)	51
Total params: 5,801		
Trainable params: 5,801		
Non-trainable params: 0		

<keras.callbacks.Historv at 0x206cee64700>

Τελικό Αποτέλεσμα (MLP):

```
# Predicting and plotting the result
trainPredict = model.predict(xtr, batch_size=batch_size)
testPredict = model.predict(xte, batch size=batch size)
# invert predictions
trainPredict = np.reshape(trainPredict, (samples*steps, features_out))
ytr2d = np.reshape(ytr, (samples*steps, features_out))
testPredict = np.reshape(testPredict, (samples*steps, features_out))
yte2d = np.reshape(yte, (samples*steps, features_out))
trainPredict = scaler.inverse_transform(trainPredict)
trainY = scaler.inverse_transform(ytr2d)
testPredict = scaler.inverse transform(testPredict)
testY = scaler.inverse_transform(yte2d)
# Calculate error
print("Test MSE: ", mean_squared_error(testY, testPredict))
print("Test MSE: ", sum(np.square(testY-testPredict))/testY.shape[0])
print( Test MAE: ", sum(abs(testY-testPredict))/testY.shape[0])
print("Test R2: ", r2_score(testY, testPredict))
print("Test R2: ", 1-(sum(np.square(testY-testPredict))/sum(np.square(testY-testY.mean()))))
Test MSE: 0.00022988309861643038
Test MSE: [0.00022988]
Test MAE: [0.01114386]
Test R2: -0.05759813323646745
Test R2: [-0.05759813]
# Finally, we check the result in a plot.
# A vertical line in a plot identifies a splitting point between
# the training and the test part.
predicted = np.concatenate((trainPredict,testPredict),axis=0)
original = np.concatenate((trainY,testY),axis=0)
predicted = np.concatenate((trainPredict,testPredict),axis=0)
index = range(0, original.shape[0])
plt.plot(index,original, 'g')
plt.plot(index,predicted, 'r')
plt.axvline(df.index[train_size], c="b")
plt.show()
 0.06
  0.04
  0.02
  0.00
-0.02
-0.04
-0.06
                100
                         200
                                  300
                                                   500
                                           400
```

Ερώτημα 2β (RNN):

Προετοιμασία για την υλοποίηση του νευρωνικού δικτύου RNN:

```
# Initialization of [samples, steps, features]
samples = train_size
steps = 1
features_in = n_in
features_out = n_out

xtr = np.reshape(xtr, (samples, steps, features_in))
ytr = np.reshape(ytr, (samples, steps, features_out))
print(xtr.shape, ytr.shape)
xte = np.reshape(xte, (samples, steps, features_in))
yte = np.reshape(yte, (samples, steps, features_out))
print(xte.shape, yte.shape)

(262, 1, 6) (262, 1, 1)
(262, 1, 6) (262, 1, 1)
```

Υλοποίηση RNN:

```
----- MLP & RNN -----
batch size = 1 #1
model = Sequential()
# MLP
# xtr, ytr = X[:train_size, :], y[:train_size]
# xte, yte = X[train_size:, :], y[train_size:]
# model.add(Dense(units=100, input_dim=xtr.shape[1], activation="relu"))
# model.add(Dense(50, activation="relu"))
# model.add(Dense(1))
# model.compile(loss='mean_squared_error', optimizer='adam')
# model.summary()
# RNN
model.add(SimpleRNN(units=100, batch_input_shape=(batch_size, xtr.shape[1],
                                                 xtr.shape[2]), activation="relu",
                                                 return_sequences=True))
model.add(Dense(50, activation="relu"))
model.add(Dense(1))
model.compile(loss='mean_squared_error', optimizer='adam')
model.summary()
model.fit(xtr,ytr, epochs=10, batch_size=batch_size, verbose=0)
4
```

Model: "sequential"

Layer (type)	Output Shape	Param #
simple_rnn (SimpleRNN)	(1, 1, 100)	10700
dense (Dense)	(1, 1, 50)	5050
dense_1 (Dense)	(1, 1, 1)	51
Total params: 15,801 Trainable params: 15,801 Non-trainable params: 0		

<keras.callbacks.History at 0x18e2d89a1f0>

Έξοδος RNN:

100

200

300

400

500

```
# Predicting and plotting the result
trainPredict = model.predict(xtr, batch_size=batch_size)
testPredict = model.predict(xte, batch_size=batch_size)
# invert predictions
trainPredict = np.reshape(trainPredict, (samples*steps, features_out))
ytr2d = np.reshape(ytr, (samples*steps, features_out))
testPredict = np.reshape(testPredict, (samples*steps, features_out))
yte2d = np.reshape(yte, (samples*steps, features_out))
trainPredict = scaler.inverse_transform(trainPredict)
trainY = scaler.inverse_transform(ytr2d)
testPredict = scaler.inverse_transform(testPredict)
testY = scaler.inverse_transform(yte2d)
# Calculate error
print("Test MSE: ", mean_squared_error(testY, testPredict))
print("Test MSE: ", sum(np.square(testY-testPredict))/testY.shape[0])
print("Test MAE: ", sum(abs(testY-testPredict))/testY.shape[0])
print("Test R2: ", r2_score(testY, testPredict))
print("Test R2: ", 1-(sum(np.square(testY-testPredict))/sum(np.square(testY-testY.mean())))))\\
Test MSE: 0.0002342335100238398
Test MSE: [0.00023423]
Test MAE: [0.01133209]
Test R2: -0.07761259715738311
Test R2: [-0.0776126]
# Finally, we check the result in a plot.
# A vertical line in a plot identifies a splitting point between
# the training and the test part.
predicted = np.concatenate((trainPredict,testPredict),axis=0)
original = np.concatenate((trainY,testY),axis=0)
predicted = np.concatenate((trainPredict,testPredict),axis=0)
index = range(0, original.shape[0])
plt.plot(index,original, 'g')
plt.plot(index,predicted, 'r')
plt.axvline(df.index[train_size], c="b")
plt.show()
 0.06
 0.04
 0.02
 0.00
-0.02
-0.04
-0.06
```