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%
% Author : Rohan Bhukar

% Load data , Examples was modified to 'examples' in my Matlab because
% of new
% version
addpath(fullfile(matlabroot,'examples','deeplearning_shared','main'));
ReadPhysionetData;

% Question 1: Part a
fprintf('Question 1: Part a \n\n')
% Describing the original dataset from Physionet
fprintf('The original counts in the dataset are: \n\n')
fprintf('Labels of classes    :   Number of Signals \n')
summary(Labels)

% Question 1: Part b
fprintf('Question 1: Part b \n\n')
% Most, but not all, of the signals in the dataset are 9000 samples in
% length. As a
% quality control measure, remove all signals that are not exactly
% 9000 samples long.
% Display the counts for each label again.

arr_sig = [];
arr_all = [];
arr_normal = [];
arr_af = [];
x = 1;
while x<=numel(Labels)
    %
    if Labels(x)=='N'
        arr_all=[arr_all; 1];
    else
        arr_all=[arr_all; 0];
    end
    %
    if numel(Signals{x})==9000
        arr_sig = [arr_sig; 1];
    else
        arr_sig = [arr_sig; 0];
    end
    x = x+1;
end

y=1;
while y<=numel(Labels)
    %
    if arr_sig(y) == 1 && arr_all(y) == 0
        arr_af = [arr_af; 1];
    else
        arr_af = [arr_af; 0];
    end
end

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        end
        %
        if arr_sig(y) == 1 && arr_all(y) == 1
            arr_normal = [arr_normal; 1];
        else
            arr_normal = [arr_normal; 0];
        end
        y = y+1;
    end

    %
    fprintf('Updates \n')
    %
    summary(Labels([find(arr_normal == 1);find(arr_af == 1)]))

% Question 1: Part c
fprintf('Question 1: Part c \n\n')
% This dataset is unbalanced, meaning the two classes are not the same
% size, which can
% introduce biases in many kinds of machine learning pipelines. Drop
% samples from
% the larger class to match the size of the smaller class. After
% balancing, randomly
% divide the data into training and validation sets with an 80:20
% ratio. Display the
% counts for each label in both the training and validation data sets.

sm_cohort_size = sum(arr_af);
random_idx = randperm(sum(arr_normal),sm_cohort_size);
j = 0;

while j == 0
    if sum(arr_af)==sum(arr_normal)
        j = 1;
    else
        val = randperm(numel(arr_normal),1);
        arr_normal(val) = 0;
    end
end

labels_mod = Labels([find(arr_normal == 1);find(arr_af == 1)]);
signals_mod = Signals([find(arr_normal == 1);find(arr_af == 1)]);

% Display the counts for each label in both the training and
% validation data sets.
split_factor = 0.8;
values = randperm(sm_cohort_size*2);

train_data =
    signals_mod(values(1:round(split_factor*sm_cohort_size*2)));

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train_labels =
    labels_mod(values(1:round(split_factor*sm_cohort_size*2)));
val_data =
    signals_mod(values(round(split_factor*sm_cohort_size*2)+1:sm_cohort_size*2));
val_labels =
    labels_mod(values(round(split_factor*sm_cohort_size*2)+1:sm_cohort_size*2));

% printing the summary here after 80:20 split
fprintf('\n\n')
summary(train_labels)
summary(val_labels)

% Question 1: Part d
fprintf('Question 1: Part d \n\n')
% create a network and train on trainign data
fprintf('\n\n')

% multiple layers for the DNN
model_lyr = [ ...
    sequenceInputLayer(1)
    bilstmLayer(100,'OutputMode','last')
    fullyConnectedLayer(2)
    softmaxLayer
    classificationLayer
]
% training options to be used from question here
options = trainingOptions('adam', ...
    'MaxEpochs',10, ...
    'MiniBatchSize', 150, ...
    'InitialLearnRate', 0.01, ...
    'SequenceLength', 1000, ...
    'GradientThreshold', 1, ...
    'ExecutionEnvironment','auto',...
    'plots','training-progress','Verbose',true);

model = trainNetwork(train_data,train_labels,model_lyr,options);

% training model on train data now

model_train_preds = classify(model,train_data,'SequenceLength',1000);

% train model accuracy, though it is displayed live on plots
train_model_accuracy = (sum(model_train_preds == train_labels)/
length(train_labels))*100
%

% Question 1: Part e
fprintf('Question 1: Part e \n\n')
% Apply your trained network to your validation data and display the
accuracy of this
% classification. Show a confusion matrix, either as a figure or text.
model_test_preds = classify(model,val_data,'SequenceLength',1000);

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test_model_accuracy = (sum(model_test_preds == val_labels)/
length(val_labels))*100

% plotting confusion matrix as asked
figure(1)
confusionchart(val_labels,model_test_preds,'ColumnSummary','column-
normalized','RowSummary','row-normalized','Title','Confusion matrix:
Testing Dataset');

% Question 2: Part a

% sub part (i)
fprintf('Question 2: Part a (i) \n\n')
[Signals,Labels] = segmentSignals(Signals,Labels);
% initial summary of data
summary(Labels)

% extracting signals and labels here
dataset_af1=Signals(Labels == 'A');
dataset_af2=Labels(Labels == 'A');
dataset_n1=Signals(Labels == 'N');
dataset_n2=Labels(Labels == 'N');

% 718 and 4937 as total sizes were chosen based on the above summary
% resulting in a ratio of approx. 1 to 7

% dividerand is used here to divide data from each class (N, A) into
train
% datasets and test datasets respectively here, but random allotment.
[af_idx1,~,af_idx2] = dividerand(718,0.8,0.0,0.2);
[n_idx1,~,n_idx2] = dividerand(4937,0.8,0.0,0.2);
af_tr1 = dataset_af1(af_idx1);
af_tr2 = dataset_af2(af_idx1);
n_tr1 = dataset_n1(n_idx1);
n_tr2 = dataset_n2(n_idx1);
af_te1 = dataset_af1(af_idx2);
af_te2 = dataset_af2(af_idx2);
n_te1 = dataset_n1(n_idx2);
n_te2 = dataset_n2(n_idx2);

% repmat for replicating(oversampling) the datasets and increasing
size to max size
train_data_x = [repmat(af_tr1(1:564),7,1); n_tr1(1:3948)];
train_data_y = [repmat(af_tr2(1:564),7,1); n_tr2(1:3948)];
test_data_x = [repmat(af_te1(1:141),7,1); n_te1(1:987)];
test_data_y = [repmat(af_te2(1:141),7,1); n_te2(1:987)];

% sub part (ii)
fprintf('Question 2: Part a (ii) \n\n')
summary(train_data_y)

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summary(test_data_y)
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% Question 2: Part b
% frequency (fs) is 300 Hz
fprintf('Question 2: Part b \n\n')
fs = 300;
% Sample input [p,f,t] =
    pspectrum(Signals{1},300,'spectrogram','TimeResolution',0.5,'Overlap',60)
% use instfreq(p,f,t) for each cell data till end for x times

% Generating features here for usefulness later
if_trf =
    cellfun(@(x)instfreq(x,fs)',train_data_x,'UniformOutput',false);
if_tef =
    cellfun(@(x)instfreq(x,fs)',test_data_x,'UniformOutput',false);
pe_trf =
    cellfun(@(x)pentropy(x,fs)',train_data_x,'UniformOutput',false);
pe_tef =
    cellfun(@(x)pentropy(x,fs)',test_data_x,'UniformOutput',false);
%instfreqTrain =
    cellfun(@(x)instfreq(pspectrum(x,fs,'spectrogram','TimeResolution',0.5,'Overlap',
%instfreqTest =
    cellfun(@(x)instfreq(pspectrum(x,fs,'spectrogram','TimeResolution',0.5,'Overlap',
%pentropyTrain =
    cellfun(@(x)pentropy(pspectrum(x,fs,'spectrogram','TimeResolution',0.5,'Overlap',
%pentropyTest =
    cellfun(@(x)pentropy(pspectrum(x,fs,'spectrogram','TimeResolution',0.5,'Overlap',
train_x_2 = cellfun(@(x,y)[x;y],if_trf,pe_trf,'UniformOutput',false);
test_x_2 = cellfun(@(x,y)[x;y],if_tef,pe_tef,'UniformOutput',false);

% computing mean and std for Z-score normalization
u_val = [train_x_2{:}];
mu = mean(u_val,2);
s_val = std(u_val,[],2);

% Z-score normalize both signals based on their mean and standard
deviation across all samples.
train_x_sd = train_x_2;
train_x_sd = cellfun(@(x)(x-mu)./
s_val,train_x_sd,'UniformOutput',false);
test_x_sd = test_x_2;
test_x_sd = cellfun(@(x)(x-mu)./
s_val,test_x_sd,'UniformOutput',false);

% plots
% Randomly choose four signals, two each of normal and AF, and plot
their
% frequency-domain features in one figure

n_sample = Signals{randperm(sm_cohort_size,1)};
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af_sample = Signals{randperm(sm_cohort_size,1)};

% extract features here
[if_a,train_a] = instfreq(af_sample,fs);
[if_n,train_n] = instfreq(n_smaple,fs);
[pe_a,train_ap] = pentropy(af_sample,fs);
[pe_n,train_np] = pentropy(n_smaple,fs);

figure(2)
subplot(2,1,1);
plot(train_n,if_n)
title('Normal Sample/signal')
xlabel('Time (in seconds)')
ylabel('Instantaneous-frequency of data')
subplot(2,1,2)
plot(train_np,pe_n)
title('Normal Sample/signal')
xlabel('Time (in seconds)')
ylabel('Spectral-entropy of data')

%
figure(3)
subplot(2,1,1)
plot(train_a,if_a)
title('Af Sample/signal')
xlabel('Time ( in seconds)')
ylabel('Instantaneous-frequency of data')
subplot(2,1,2)
plot(train_ap,pe_a)
title('Af Sample/signal')
xlabel('Time (in seconds)')
ylabel('Spectral-entropy of data')

% Question 2: Part c
%
fprintf('Question 2: Part c \n\n')

% different layers as requested above
% Input layer
% biLSTM layer with asked output mode
% full connected layer which takes input from biLSTM, also 2 for
specifying
% 2 classes here
% a softmax layer to normalize the output of a network to a
probability distribution over predicted output classes
% a classification layer computes the cross-entropy loss for
classification and weighted classification tasks with mutually
exclusive classes

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modl_layers = [ ...
    sequenceInputLayer(2)
    bilstmLayer(100,'OutputMode','last')
    fullyConnectedLayer(2)
    softmaxLayer
    classificationLayer
];
% options to be used in model training
options = trainingOptions('adam','MaxEpochs',30, 'MiniBatchSize',
    150, ...
    'InitialLearnRate', 0.01, 'GradientThreshold',
    1, 'ExecutionEnvironment', ...
    "auto",'plots','training-progress','Verbose',true);

% train network created above on training data
model2 = trainNetwork(train_x_sd,train_data_y,modl_layers,options);
model_train_preds2 = classify(model2,train_x_sd);
train_model_accuracy = sum(model_train_preds2 == train_data_y)/
length(train_data_y)*100

% apply train network to val data
fprintf('Question 2: Part d \n\n')
model_test_preds2 = classify(model2,test_x_sd);
test_model_accuracy = sum(model_test_preds2 == test_data_y)/
length(test_data_y)*100

% confusion matrix for test data as asked

figure(4)
confusionchart(test_data_y,model_test_preds2,'ColumnSummary','column-
normalized','RowSummary','row-normalized','Title','Confusion matrix :
Testing dataset');

% Question 3 :
fprintf('Question 3: \n\n')
% Results respectively have been shown in the plots and data files,
% and outputs generated while running the algo.

% From my current understanding, the approach number = 2, has
% performed
% better in this case. The method / approach here in case 2 , had
% higher
% performance because in this case the useful features were extracted
% from
% this dataset prior before we started implementing the deep-learning
% algorithm of LSTMs.
% While in this 2nd case the step we performed for extracting the 2
% time-series signals based on their freq domain features (spectral
% entropy

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% and instantenous frequency), both of which have helped in incerasing
the
% training accuracy. They also have resulted in reduced training time
due
% to shorter lengths of sequences.
% What we learned here today, is that the feature extraction steps are
% important steps when working with deep learning algorithms such as
LSTMs,
% and for problems like this in the future even with different sets of
data
% we should try to attempt to apply the above feature extraction
steps.
```

Question 1: Part a

The original counts in the dataset are:

```
Labels of classes    : Number of Signals
      A             738
      N             5050
```

Question 1: Part b

```
Updates
      A             499
      N             3678
```

Question 1: Part c

```
      A             397
      N             401
      A             102
      N              98
```

Question 1: Part d

```
model_lyr =
```

5x1 Layer array with layers:

```
1  ''  Sequence Input           Sequence input with 1 dimensions
2  ''  BiLSTM                   BiLSTM with 100 hidden units
3  ''  Fully Connected          2 fully connected layer
4  ''  Softmax                  softmax
5  ''  Classification Output    crossentropyex
```

Training on single CPU.

```
|
=====
| Epoch | Iteration | Time Elapsed | Mini-batch | Mini-batch
| Base Learning |
|           |           | (hh:mm:ss)  | Accuracy  | Loss
|           |           |              |           |
|           |           |              |           |
|           |           |              |           |
```



```

/
=====
/      1 /      1 /      00:00:33 /      58.00% /      0.7296
/      0.0100 /
/      1 /      10 /      00:03:04 /      44.67% /      0.7235
/      0.0100 /
/
=====

```

`train_model_accuracy =`

53.1328

Question 1: Part e

`test_model_accuracy =`

46

Question 2: Part a (i)

A 718

N 4937

Question 2: Part a (ii)

A 3948

N 3948

A 987

N 987

Question 2: Part b

Question 2: Part c

Training on single CPU.

```

/
=====
/ Epoch / Iteration / Time Elapsed / Mini-batch / Mini-batch
/ Base Learning /
/      /      / (hh:mm:ss) / Accuracy / Loss
/      Rate      /
/
=====
/      1 /      1 /      00:00:08 /      58.67% /      0.6883
/      0.0100 /
/      1 /      50 /      00:01:32 /      67.33% /      0.5686
/      0.0100 /
/      2 /      100 /      00:03:05 /      74.00% /      0.5538
/      0.0100 /
/      3 /      130 /      00:03:59 /      70.67% /      0.6657
/      0.0100 /
/
=====

```

`train_model_accuracy =`

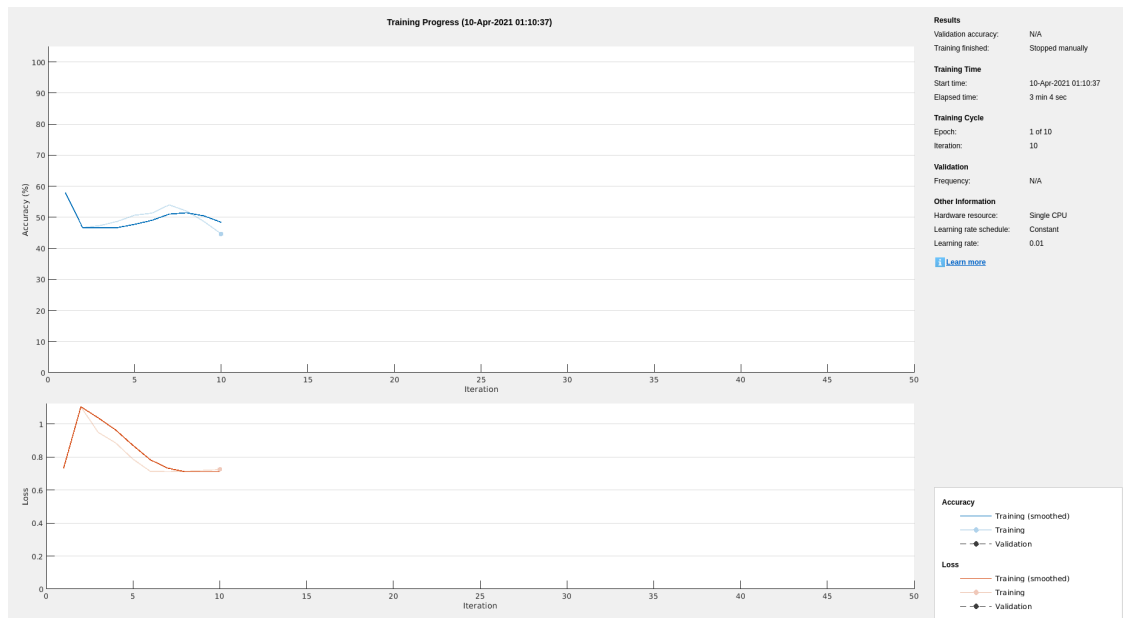
`75.0127`

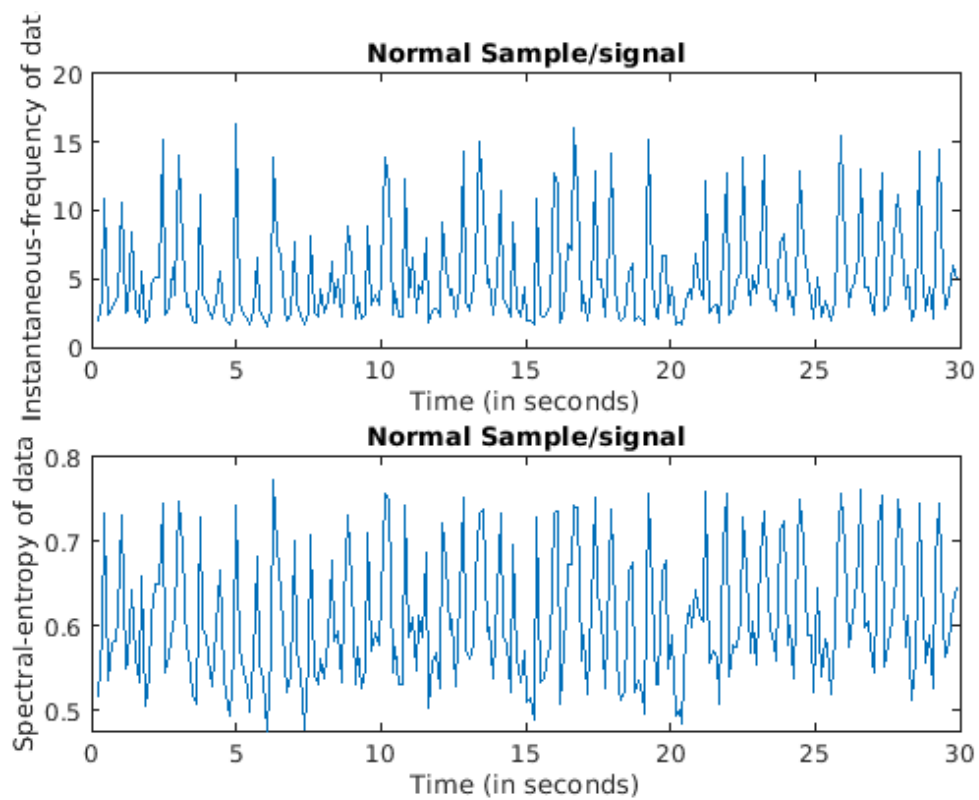
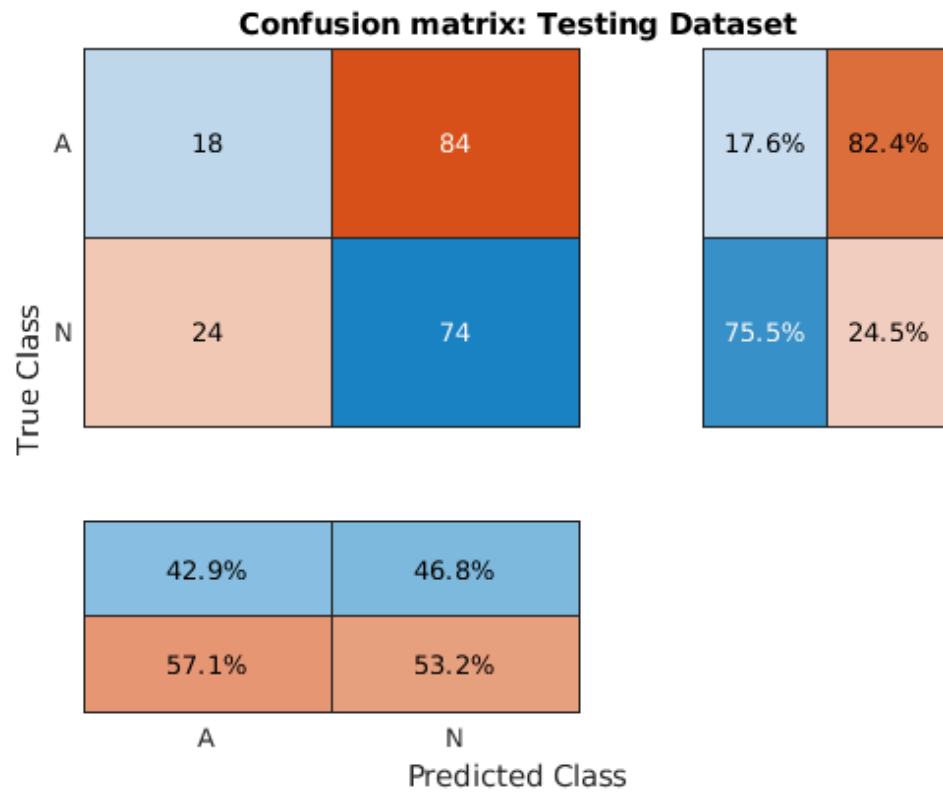
*Question 2: Part d*

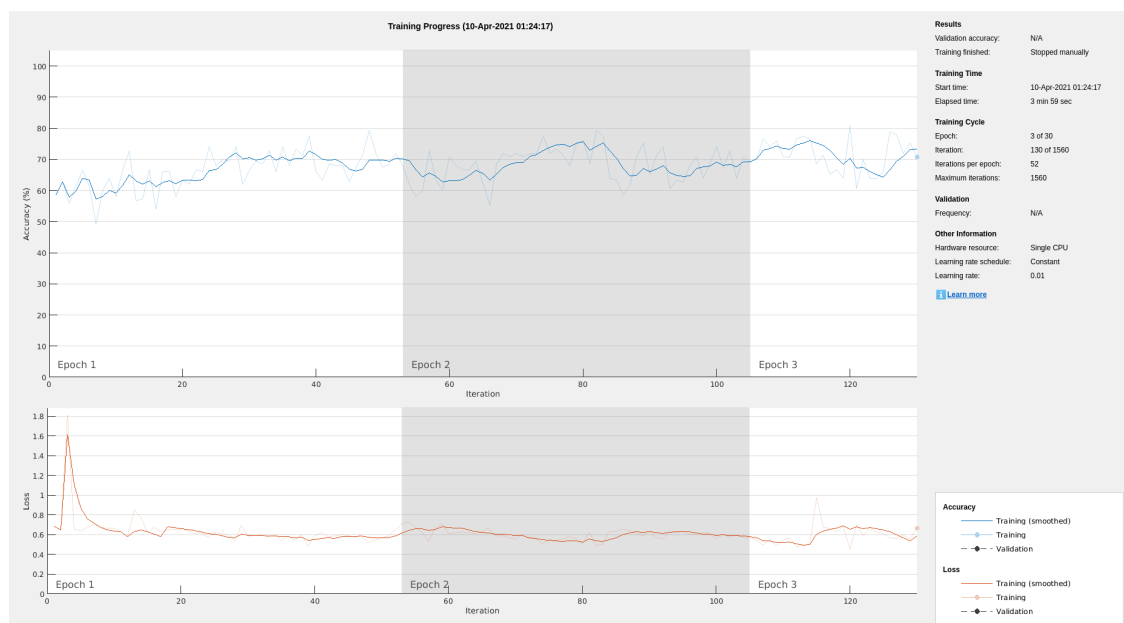
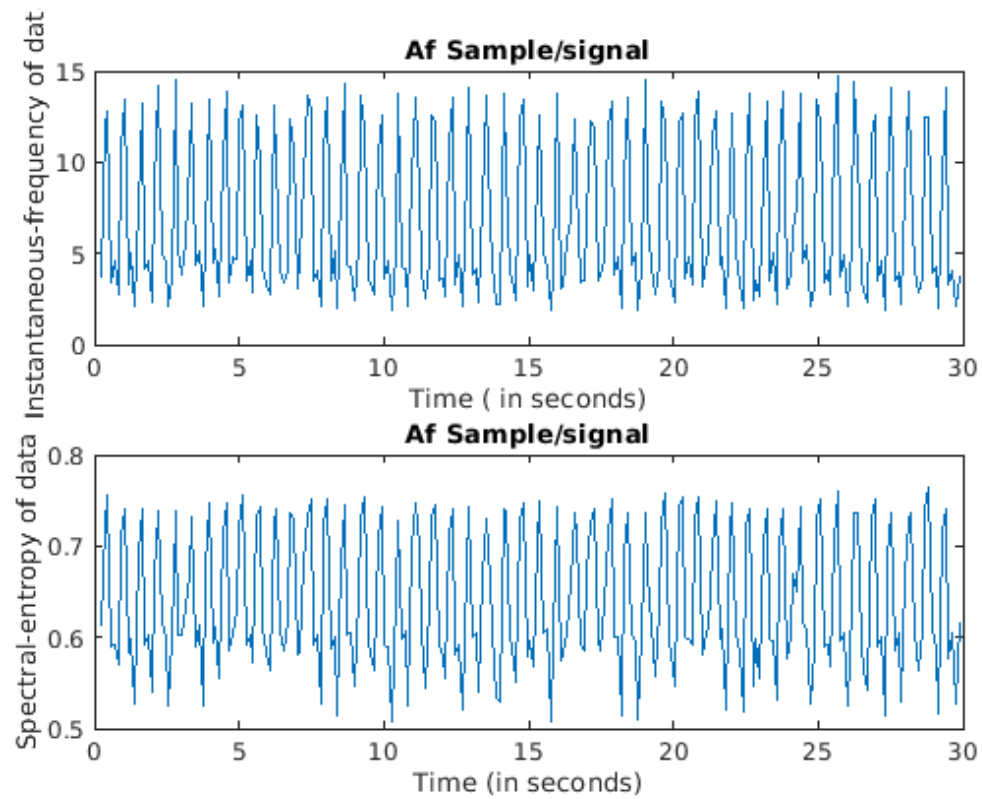
`test_model_accuracy =`

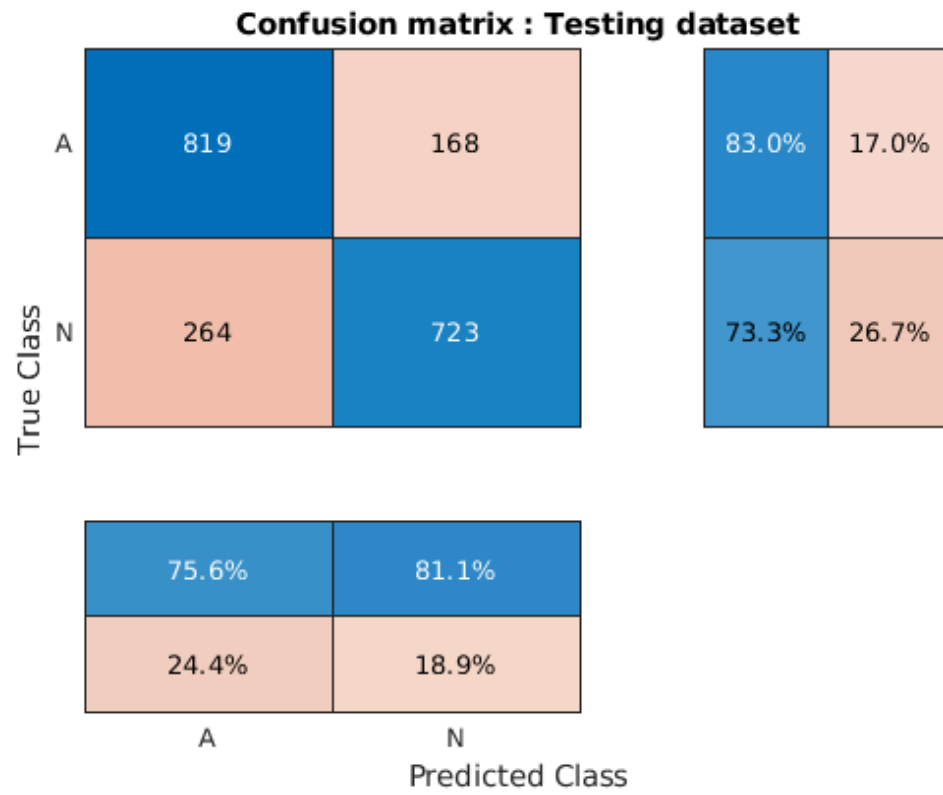
`78.1155`

*Question 3:*









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