

HAR_LSTM-MODEL

June 29, 2019

```
In [1]: # Importing Libraries
import pandas as pd
import numpy as np
import os
import pickle

# Import Model related packages
import tensorflow as tf
from keras import backend as K
from keras.models import Sequential
from keras.layers import LSTM
from keras.layers.core import Dense, Dropout
from keras.callbacks import EarlyStopping, ModelCheckpoint

# Model evaluation related packages
from sklearn.metrics import confusion_matrix
from sklearn.metrics import precision_recall_fscore_support

# for one hot encoding of labels
from sklearn.preprocessing import OneHotEncoder

# Plot related packages
import matplotlib.pyplot as plt
import seaborn as sns
sns.set()

# Presenting the results
from prettytable import PrettyTable
```

Using TensorFlow backend.

1 HumanActivityRecognition

This project is to build a model that predicts the human activities such as Walking, Walking_Upstairs, Walking_Downstairs, Sitting, Standing or Laying.

This dataset is collected from 30 persons(referred as subjects in this dataset), performing different activities with a smartphone to their waists. The data is recorded with the help of sensors

(accelerometer and Gyroscope) in that smartphone. This experiment was video recorded to label the data manually.

1.1 How data was recorded

By using the sensors(Gyroscope and accelerometer) in a smartphone, they have captured '3-axial linear acceleration'($tAcc-XYZ$) from accelerometer and '3-axial angular velocity' ($tGyro-XYZ$) from Gyroscope with several variations.

prefix ' t ' in those metrics denotes time.

suffix ' XYZ ' represents 3-axial signals in X , Y, and Z directions.

1.1.1 Feature names

1. These sensor signals are preprocessed by applying noise filters and then sampled in fixed-width windows(sliding windows) of 2.56 seconds each with 50% overlap. ie., each window has 128 readings.
2. From Each window, a feature vector was obtained by calculating variables from the time and frequency domain. > In our dataset, each datapoint represents a window with different readings
3. The acceleration signal was separated into Body and Gravity acceleration signals($tBodyAcc-XYZ$ and $tGravityAcc-XYZ$) using some low pass filter with corner frequency of 0.3Hz.
4. After that, the body linear acceleration and angular velocity were derived in time to obtain *jerk signals* ($tBodyAccJerk-XYZ$ and $tBodyGyroJerk-XYZ$).
5. The magnitude of these 3-dimensional signals were calculated using the Euclidian norm. These magnitudes are represented as features with names like $tBodyAccMag$, $tGravityAccMag$, $tBodyAccJerkMag$, $tBodyGyroMag$ and $tBodyGyroJerkMag$.
6. Finally, We've got frequency domain signals from some of the available signals by applying a FFT (Fast Fourier Transform). These signals obtained were labeled with *prefix ' f '* just like original signals with *prefix ' t '*. These signals are labeled as $fBodyAcc-XYZ$, $fBodyGyroMag$ etc.,.
7. These are the signals that we got so far.
 - $tBodyAcc-XYZ$
 - $tGravityAcc-XYZ$
 - $tBodyAccJerk-XYZ$
 - $tBodyGyro-XYZ$
 - $tBodyGyroJerk-XYZ$
 - $tBodyAccMag$
 - $tGravityAccMag$
 - $tBodyAccJerkMag$
 - $tBodyGyroMag$
 - $tBodyGyroJerkMag$
 - $fBodyAcc-XYZ$

- fBodyAccJerk-XYZ
 - fBodyGyro-XYZ
 - fBodyAccMag
 - fBodyAccJerkMag
 - fBodyGyroMag
 - fBodyGyroJerkMag
8. We can estimate some set of variables from the above signals. ie., We will estimate the following properties on each and every signal that we recorded so far.
- *mean()*: Mean value
 - *std()*: Standard deviation
 - *mad()*: Median absolute deviation
 - *max()*: Largest value in array
 - *min()*: Smallest value in array
 - *sma()*: Signal magnitude area
 - *energy()*: Energy measure. Sum of the squares divided by the number of values.
 - *iqr()*: Interquartile range
 - *entropy()*: Signal entropy
 - *arCoeff()*: Autoregression coefficients with Burg order equal to 4
 - *correlation()*: correlation coefficient between two signals
 - *maxInds()*: index of the frequency component with largest magnitude
 - *meanFreq()*: Weighted average of the frequency components to obtain a mean frequency
 - *skewness()*: skewness of the frequency domain signal
 - *kurtosis()*: kurtosis of the frequency domain signal
 - *bandsEnergy()*: Energy of a frequency interval within the 64 bins of the FFT of each window.
 - *angle()*: Angle between two vectors.
9. We can obtain some other vectors by taking the average of signals in a single window sample. These are used on the angle() variable''
- gravityMean
 - tBodyAccMean
 - tBodyAccJerkMean
 - tBodyGyroMean
 - tBodyGyroJerkMean

1.1.2 Y_Labels(Encoded)

- In the dataset, Y_labels are represented as numbers from 1 to 6 as their identifiers.
 - WALKING as 1
 - WALKING_UPSTAIRS as 2
 - WALKING_DOWNSTAIRS as 3
 - SITTING as 4
 - STANDING as 5
 - LAYING as 6

1.2 Train and test data were saperated

- The readings from **70%** of the volunteers were taken as *trianing data* and remaining **30%** subjects recordings were taken for *test data*

1.3 Data

- All the data is present in 'UCI_HAR_dataset/' folder in present working directory.
 - Feature names are present in 'UCI_HAR_dataset/features.txt'
 - *Train Data*
 - * 'UCI_HAR_dataset/train/X_train.txt'
 - * 'UCI_HAR_dataset/train/subject_train.txt'
 - * 'UCI_HAR_dataset/train/y_train.txt'
 - *Test Data*
 - * 'UCI_HAR_dataset/test/X_test.txt'
 - * 'UCI_HAR_dataset/test/subject_test.txt'
 - * 'UCI_HAR_dataset/test/y_test.txt'

1.4 Data Size :

27 MB

2 Quick overview of the dataset :

- Accelerometer and Gyroscope readings are taken from 30 volunteers(referred as subjects) while performing the following 6 Activities.
 1. Walking
 2. WalkingUpstairs
 3. WalkingDownstairs
 4. Standing
 5. Sitting
 6. Lying.
- Readings are divided into a window of 2.56 seconds with 50% overlapping.
- Accelerometer readings are divided into gravity acceleration and body acceleration readings, which has x,y and z components each.
- Gyroscope readings are the measure of angular velocities which has x,y and z components.
- Jerk signals are calculated for BodyAcceleration readings.
- Fourier Transforms are made on the above time readings to obtain frequency readings.
- Now, on all the base signal readings., mean, max, mad, sma, arcoefficient, engery-bands,entropy etc., are calculated for each window.

- We get a feature vector of 561 features and these features are given in the dataset.
- Each window of readings is a datapoint of 561 features.

2.1 Problem Framework

- 30 subjects(volunteers) data is randomly split to 70%(21) test and 30%(7) train data.
- Each datapoint corresponds one of the 6 Activities.

3 Configs

```
In [2]: # base directory for dataset
data_base_dir = '/home/nisheel-s/RUN_AMT/DS/CS11_HUMAN_ACTIVITY_RECOGNITION/'

# classification related configs
num_classes = 6
class_names_list = ['WALKING', 'WALKING_UPSTAIRS', 'WALKING_DOWNSTAIRS',
                    'STANDING', 'SITTING', 'LAYING']
class_labels_list = list(range(1, num_classes + 1))

# dataset size related configs
dataset_size = -1 # set -1 for using full size
```

4 UTIL functions

```
In [3]: def get_confusion_matrix(actual_list, predicted_list, title_suffix=str(), plot_fig=True)
        """
        This function plots the confusion matrix given ground truth and predicted
        """

        conf_matrix = confusion_matrix(actual_list, predicted_list,
                                       labels=list(range(0, num_classes)))

        conf_df = pd.DataFrame(conf_matrix, columns=class_names_list)
        conf_df.index = class_names_list

        # plot confusion matrix
        if plot_fig:
            plt.figure(figsize=(9,6))
            sns.heatmap(conf_df, annot=True, annot_kws={'size': 8}, fmt='g', cmap='YlGnBu',
                        cbar_kws={'label': 'prediction_count', 'format': '%d'})
            plt.yticks(rotation=0)
            plt.xticks(rotation=90)
            plt.xlabel('Predicted')
            plt.ylabel('Actual')
            plt.title('Confusion Matrix -' + title_suffix)
            plt.show()
```

```

    return conf_df

In [4]: def get_precision_recall_matrix(conf_matrix, title_suffix=str(), plot_fig=True):

    # compute precision matrix
    precision_matrix = conf_matrix.div(conf_matrix.sum(axis=0), axis=1) * 100.0

    # compute recall matrix
    recall_matrix = conf_matrix.div(conf_matrix.sum(axis=1), axis=0) * 100.0

    # plot both the matrices
    if plot_fig:
        # plot precision matrix
        plt.figure(figsize=(9,6))
        sns.heatmap(precision_matrix, annot=True, annot_kws={'size': 8},
                    fmt='.4f', cmap='YlGnBu',
                    cbar_kws={'label': 'percentage', 'format': '%.2f'})
        plt.yticks(rotation=0)
        plt.xticks(rotation=90)
        plt.xlabel('Predicted')
        plt.ylabel('Actual')
        plt.title('Precision Matrix -' + title_suffix)
        plt.show()

        # plot recall matrix
        plt.figure(figsize=(9,6))
        sns.heatmap(recall_matrix, annot=True, annot_kws={'size': 8},
                    fmt='.4f', cmap='YlGnBu',
                    cbar_kws={'label': 'percentage', 'format': '%.2f'})
        plt.yticks(rotation=0)
        plt.xticks(rotation=90)
        plt.xlabel('Predicted')
        plt.ylabel('Actual')
        plt.title('Recall Matrix -' + title_suffix)
        plt.show()

    # return as a tuple
    return (precision_matrix, recall_matrix,)

In [5]: def get_classification_report(actual, predicted, title_suffix=str(), plot_fig=True):

    # compute performance df
    eval_matrix = precision_recall_fscore_support(actual, predicted,
                                                  labels=list(range(0, num_classes)))

    eval_df = pd.DataFrame(list(eval_matrix), columns=list(range(0, num_classes)))
    eval_df.index = ['Precision', 'Recall', 'Fscore', 'Support']

```

```

eval_df.columns = class_names_list

# normalize the performace df
eval_df_normed = eval_df * 100.0
eval_df_normed.loc['Support', class_names_list] /= eval_df_normed.loc['Support',
                                                                    class_names_list].sum()

eval_df_normed.iloc[3:4, :] *= 100.0

# plot the classification report
if plot_fig:
    plt.figure(figsize=(9,6))
    sns.heatmap(eval_df_normed, annot=True, annot_kws={'size': 8},
                fmt='.4f', cmap='YlGnBu',
                cbar_kws={'label': 'Percentage', 'format': '%.2f'})
    plt.yticks(rotation=0)
    plt.xticks(rotation=90)

    plt.xlabel('Classes')
    plt.ylabel('Metrics')
    plt.title('Classification Report on ' + title_suffix + ' Data')
    plt.show()

return eval_df_normed

```

```

In [6]: def plot_loss_curve(hist_obj):
        """
        This function plots the loss curve
        """

        # get train , validation loss information from object
        train_loss_list = list(hist_obj['loss'])
        validation_loss_list = list(hist_obj['val_loss'])

        # get x-label list
        epoch_list = range(1, len(train_loss_list) + 1 )

        # plot the performace
        # plot both train, validation curve
        plt.plot(epoch_list, train_loss_list,
                 label='Train Loss', color='r')
        plt.plot(epoch_list, validation_loss_list,
                 label='Validation Loss', color='b')
        plt.xlabel('Training Epoch')
        plt.ylabel('Multiclass Log Loss')
        plt.title('Training Loss Vs Validation Loss')
        plt.legend()
        plt.show()

```

```

In [7]: def plot_accuracy_curve(hist_obj):
        """
        This is a helper function for plotting accuracy
        """

        # extract train, validation accuracy metrics
        train_acc_list = list(hist_obj['acc'])
        val_acc_list = list(hist_obj['val_acc'])

        # get x-label list
        epoch_list = range(1, len(train_acc_list) + 1 )

        # plot both train, validation curve
        plt.plot(epoch_list, train_acc_list, label='Train Accuracy', color='r')
        plt.plot(epoch_list, val_acc_list, label='Validation Accuracy', color='b')
        plt.xlabel('Training Epoch')
        plt.ylabel('Accuracy')
        plt.title('Training Accuracy Vs Validation Accuracy')
        plt.legend()
        plt.show()

```

5 Data

```

In [8]: # declare one hot encoder for label
        one_hot_encoder = OneHotEncoder(sparse=False, categories='auto')
        one_hot_encoder.fit(np.array(class_labels_list).reshape(num_classes, 1))

```

```

Out[8]: OneHotEncoder(categorical_features=None, categories='auto',
                        dtype=<class 'numpy.float64'>, handle_unknown='error',
                        n_values=None, sparse=False)

```

```

In [9]: # Utility function to load the features
        def load_features(data_base_dir, dataset_type):

            # declare all the signals in a list
            SIGNALS = [
                'body_acc_x',
                'body_acc_y',
                'body_acc_z',
                'body_gyro_x',
                'body_gyro_y',
                'body_gyro_z',
                'total_acc_x',
                'total_acc_y',
                'total_acc_z' ]

            # declare a list for holding the entire data frame

```



```

signals_data = list()

# do for all signals in the list
for signal in SIGNALS:

    # set the file path
    file_path = os.path.join(data_base_dir,
                              f'{dataset_type}/Inertial Signals/{signal}_{dataset_type}.csv')

    # read the file
    tdf = pd.read_csv(file_path, delim_whitespace=True, header=None)
    # update the list
    signals_data.append(tdf.values)

# Transpose is used to change the dimensionality of the output,
# aggregating the signals by combination of sample/timestep.
# Resultant shape is (7352 train/2947 test samples, 128 timesteps, 9 signals)
signals_data = np.transpose(signals_data, (1, 2, 0))

return signals_data

```

```

In [10]: def load_labels(data_base_dir, dataset_type):

    # set the file path
    file_path = os.path.join(data_base_dir, f'{dataset_type}/y_{dataset_type}.txt')
    # read the file
    tdf = pd.read_csv(file_path, delim_whitespace=True, header=None, squeeze=True)

    print('Distribution of labels :\n', tdf.value_counts())

    # get labels in one-hot encoded representation
    labels = one_hot_encoder.transform(tdf.values.reshape(len(tdf), 1)).astype(int)

    return labels

```

5.1 Raw data signals Information

Signals are from Accelerometer and Gyroscope

The signals are in x,y,z directions

Triaxial acceleration from the accelerometer is total acceleration.

Sensor signals are filtered to have only body acceleration
excluding the acceleration due to gravity

```

In [11]: # Load the features
X_train = load_features(data_base_dir, 'train')
X_test = load_features(data_base_dir, 'test')
# Load the labels
y_train = load_labels(data_base_dir, 'train')

```

```
y_test = load_labels(data_base_dir, 'test')
```

```
if dataset_size > 0:  
    X_train = X_train[0:dataset_size]  
    X_test = X_test[0:dataset_size]  
    y_train = y_train[0:dataset_size]  
    y_test = y_test[0:dataset_size]
```

Distribution of labels :

```
6    1407  
5    1374  
4    1286  
1    1226  
2    1073  
3     986
```

Name: 0, dtype: int64

Distribution of labels :

```
6     537  
5     532  
1     496  
4     491  
2     471  
3     420
```

Name: 0, dtype: int64

```
In [12]: print('Train features shape : ', X_train.shape)  
         print('Train labels shape : ', y_train.shape)  
         print('Test features shape : ', X_test.shape)  
         print('Test labels shape : ', y_test.shape)
```

Train features shape : (7352, 128, 9)

Train labels shape : (7352, 6)

Test features shape : (2947, 128, 9)

Test labels shape : (2947, 6)

6 MODEL

```
In [13]: def build_model_a(timesteps, input_dim, h_params):
```

```
    # Initiliazing the sequential model  
    model = Sequential()
```

```
    # extract the hyper params  
    n_hidden = h_params[0]  
    dr_rate = h_params[1]
```

```

# Configuring the parameters
model.add(LSTM(n_hidden, input_shape=(timesteps, input_dim)))

# Adding a dropout layer
model.add(Dropout(dr_rate))

# Adding a dense output layer with sigmoid activation
model.add(Dense(num_clasess, activation='sigmoid'))

print(model.summary())

return model

```

```
In [14]: def build_model_b(timesteps, input_dim, h_params):
```

```

# Initiliazing the sequential model
model = Sequential()

# Extract Hyperparameters

# 1 - extract the number of neurons
n_hidden_1 = h_params[0]
n_hidden_2 = h_params[1]

# 2 - extract dropout rates
dr_rate_1 = h_params[2]
dr_rate_2 = h_params[3]

# Configuring the parameters
model.add(LSTM(n_hidden_1, input_shape=(timesteps, input_dim),
              return_sequences=True))
model.add(Dropout(dr_rate_1))
model.add(LSTM(n_hidden_2, return_sequences=False))

# Adding a dropout layer
model.add(Dropout(dr_rate_2))

# Adding a dense output layer with sigmoid activation
model.add(Dense(num_clasess, activation='sigmoid'))

print(model.summary())

return model

```

```
In [15]: def train_model(model, model_name, X_train, y_train, num_epochs):
```

```

# Compiling the model
model.compile(loss='categorical_crossentropy',

```

```

optimizer='adam',
metrics=['accuracy'])

# monitor the performace of model on every epoch
checkpoint = ModelCheckpoint('./model/' + model_name, monitor='val_loss',
                             verbose=0, save_best_only=True, mode='auto')

# add early stopping
early_stop = EarlyStopping(monitor='val_loss', min_delta=0,
                             patience=5, verbose=0, mode='auto')

callbacks_list = [checkpoint, early_stop]

# Train the model
hist_obj = model.fit(X_train, y_train, epochs=num_epochs, batch_size=batch_size,
                     validation_data=(X_test, y_test),
                     callbacks=callbacks_list, verbose=0)
hist_obj = hist_obj.history

# write train history to a file
hist_path = './results/' + model_name[0:-3] + '_train_history.pkl'
pickle_out = open(hist_path, 'wb')
pickle.dump(hist_obj, pickle_out)
pickle_out.close()

return hist_obj

```

```

In [16]: def evaluate_model(model, model_name, X, y, title_suffix=str(), plot_fig=True):

```

```

    # load weights from the saved model
    model_path = './model/' + model_name
    model.load_weights(model_path)

    # Compile model (required to make predictions)
    model.compile(loss='categorical_crossentropy', optimizer='adam',
                  metrics=['accuracy'])
    print('Restored best model weights from saved file', model_path)

    # evaluate model
    score = model.evaluate(X, y)

    # predict using model & get the labels
    predicted_labels = model.predict(X)
    predicted_labels_list = predicted_labels.argmax(axis=1)

    # format the actual list

```

```

y_list = y.argmax(axis=1)

# 1. Create confusion matrix
conf_df = get_confusion_matrix(y_list, predicted_labels_list,
                               title_suffix, plot_fig)

# 2. Precision, Recall matrix
pre_matrix, rec_matrix = get_precision_recall_matrix(conf_df,
                                                    title_suffix,
                                                    plot_fig)

# 3. Evaluation metrics
eval_df = get_classification_report(y_list, predicted_labels_list,
                                   title_suffix, plot_fig)

# convert accuracy to percentage
score[1] *= 100.0

# extract individual fields
return score

```

7 Run Models

```

In [17]: # Initializing parameters
num_epochs = 30
batch_size = 16

timesteps = len(X_train[0])
input_dim = len(X_train[0][0])

print('Number of classes :', num_classes)
print('Number of timesteps :', timesteps)
print('Input data dimension :', input_dim)

```

```

Number of classes : 6
Number of timesteps : 128
Input data dimension : 9

```

```

In [18]: # Importing tensorflow
np.random.seed(42)
tf.set_random_seed(42)

```

```

In [19]: # Configuring a session
session_conf = tf.ConfigProto(
    intra_op_parallelism_threads=1,
    inter_op_parallelism_threads=1
)

```

```
In [20]: sess = tf.Session(graph=tf.get_default_graph(), config=session_conf)
        K.set_session(sess)
```

7.1 a) Single-Layered Architecture

7.1.1 Architecture 1

```
In [21]: # Build the Model
        h_params_a1 = (32, 0.32,)
        model_a1 = build_model_a(timesteps, input_dim, h_params_a1)
```

WARNING:tensorflow:From /home/nisheel-s/anaconda3/lib/python3.6/site-packages/keras/backend/tensorflow_backend.py:1445: Instructions for updating:
keep_dims is deprecated, use keepdims instead

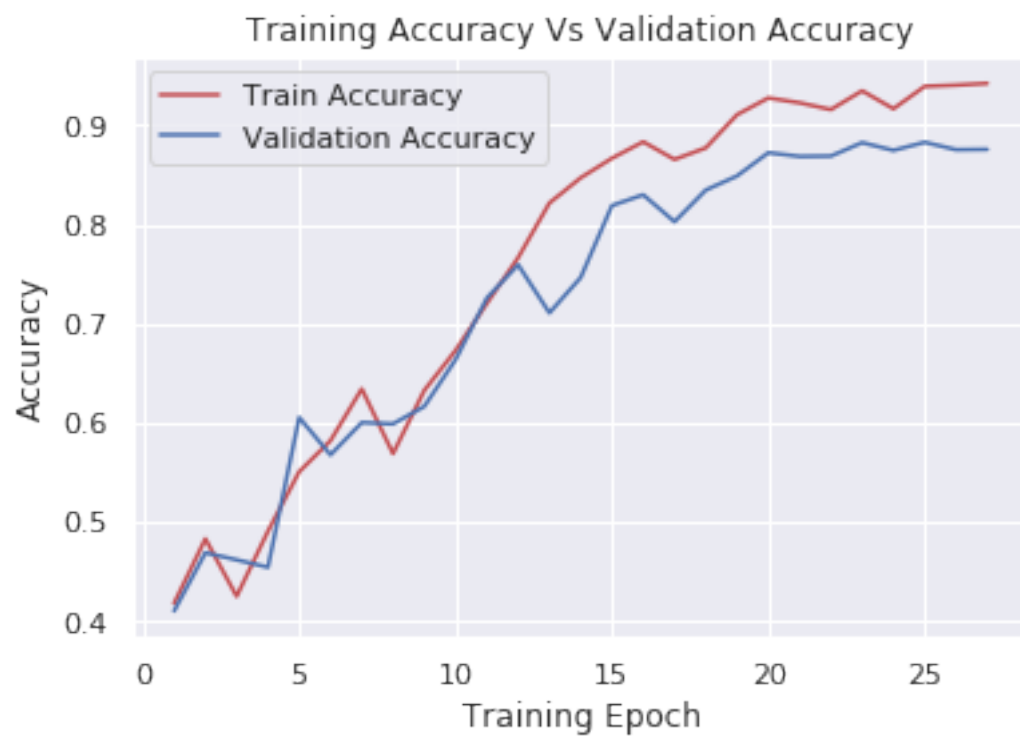
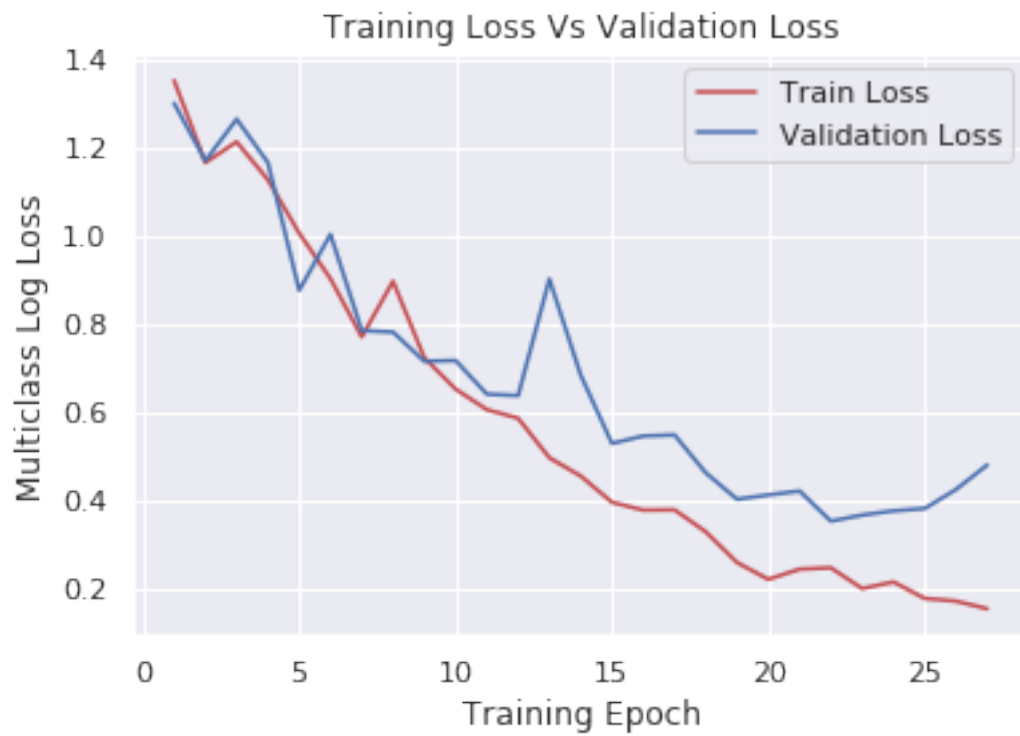
Layer (type)	Output Shape	Param #
lstm_1 (LSTM)	(None, 32)	5376
dropout_1 (Dropout)	(None, 32)	0
dense_1 (Dense)	(None, 6)	198

Total params: 5,574
 Trainable params: 5,574
 Non-trainable params: 0

```
In [22]: # Train the Model
        model_a1_name = 'Single_Layered_LSTM_1.h5'
        hist_model_a1 = train_model(model_a1, model_a1_name, X_train, y_train, num_epochs)
```

WARNING:tensorflow:From /home/nisheel-s/anaconda3/lib/python3.6/site-packages/keras/backend/tensorflow_backend.py:1445: Instructions for updating:
keep_dims is deprecated, use keepdims instead

```
In [23]: # plot loss, accuracy curves
        plot_loss_curve(hist_model_a1)
        plot_accuracy_curve(hist_model_a1)
```

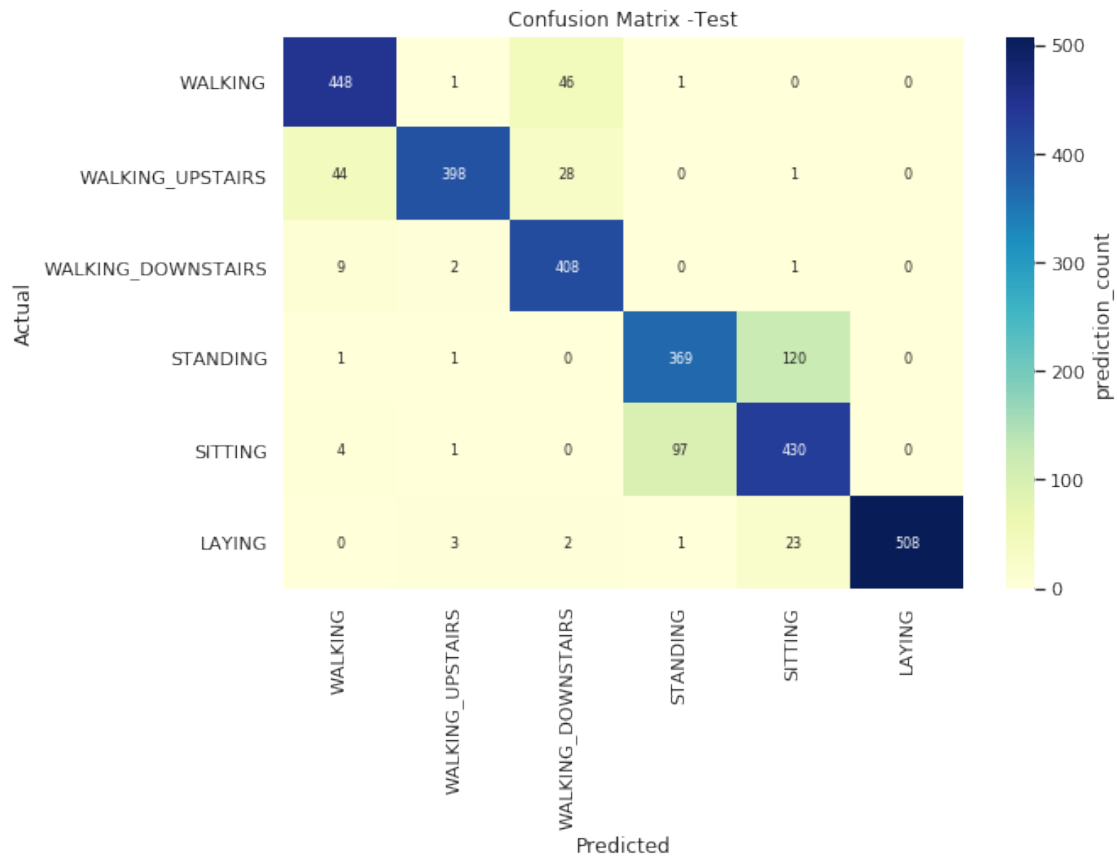


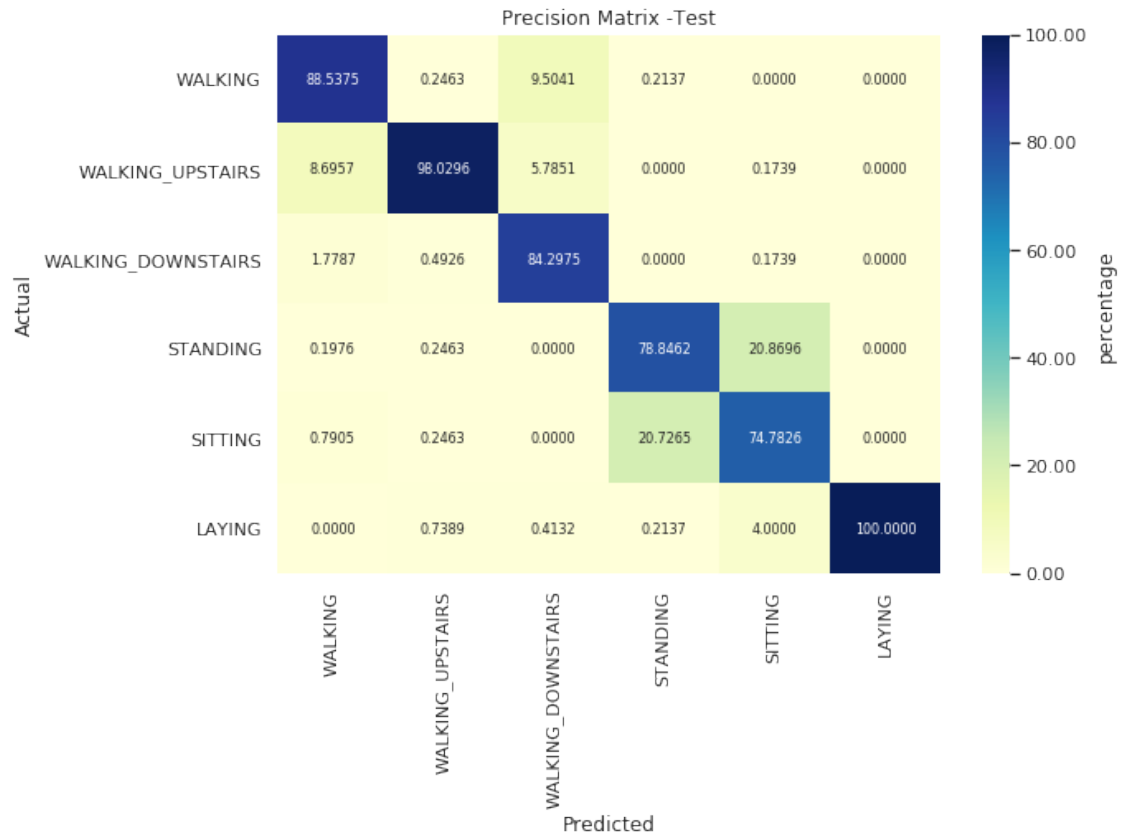
```
In [24]: # Evaluate the model on Test data
train_score_model_a1 = evaluate_model(model_a1, model_a1_name, X_train, y_train,
                                     title_suffix='Train', plot_fig=False)
train_score_model_a1 = [round(item, 4) for item in train_score_model_a1]
```

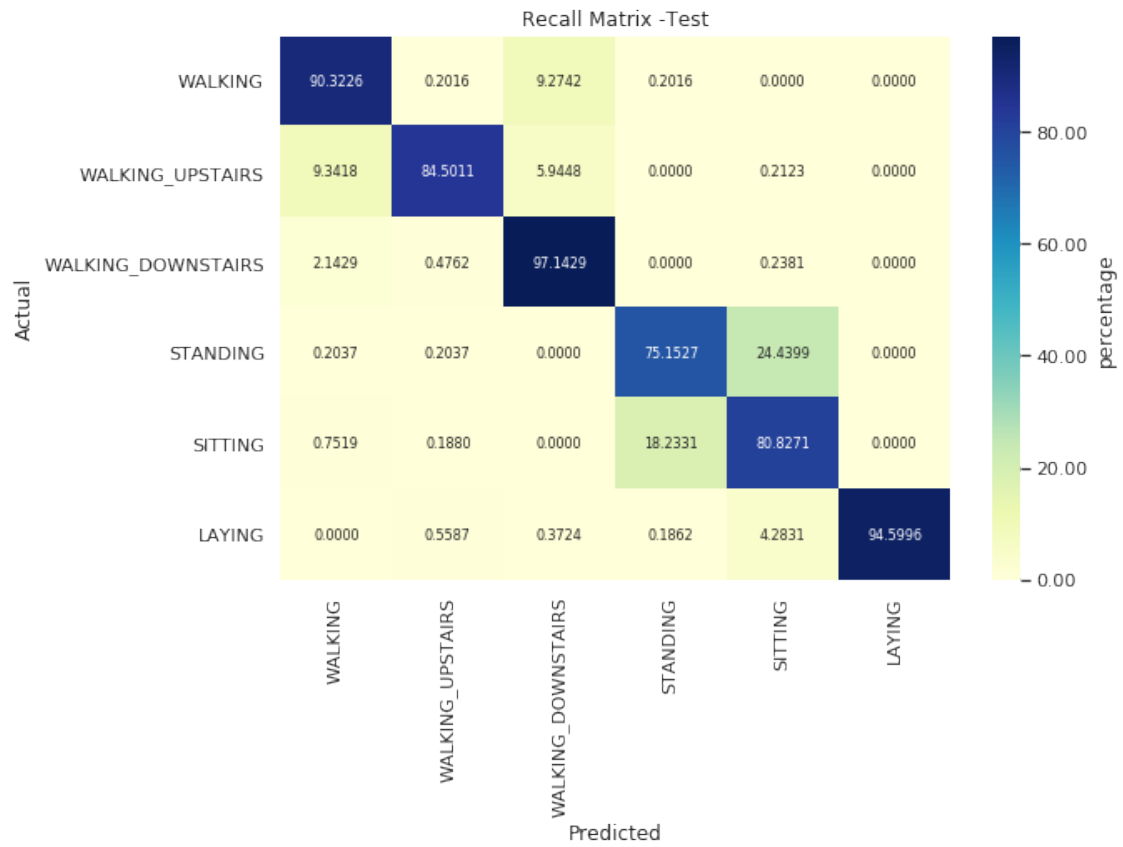
Restored best model weights from saved file ./model/Single_Layered_LSTM_1.h5
7352/7352 [=====] - 2s 304us/step

```
In [25]: # Evaluate the model on Test data
test_score_model_a1 = evaluate_model(model_a1, model_a1_name, X_test, y_test,
                                    title_suffix='Test', plot_fig=True)
test_score_model_a1 = [round(item, 4) for item in test_score_model_a1]
```

Restored best model weights from saved file ./model/Single_Layered_LSTM_1.h5
2947/2947 [=====] - 1s 351us/step









```
In [26]: table_entry_model_a1 = ['1 Layered LSTM(1)', h_params_a1] + train_score_model_a1 + \
      test_score_model_a1
```

7.1.2 Architecture 2

```
In [27]: # Build the Model
      h_params_a2 = (48, 0.40,)
      model_a2 = build_model_a(timesteps, input_dim, h_params_a2)
```

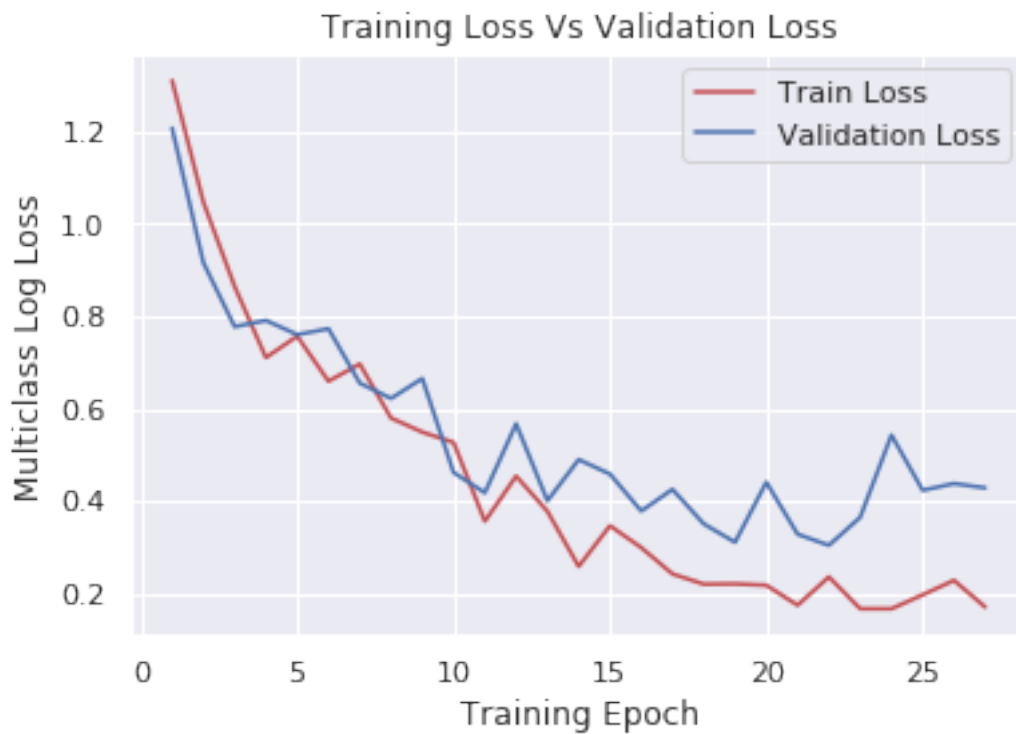
Layer (type)	Output Shape	Param #
lstm_2 (LSTM)	(None, 48)	11136
dropout_2 (Dropout)	(None, 48)	0
dense_2 (Dense)	(None, 6)	294

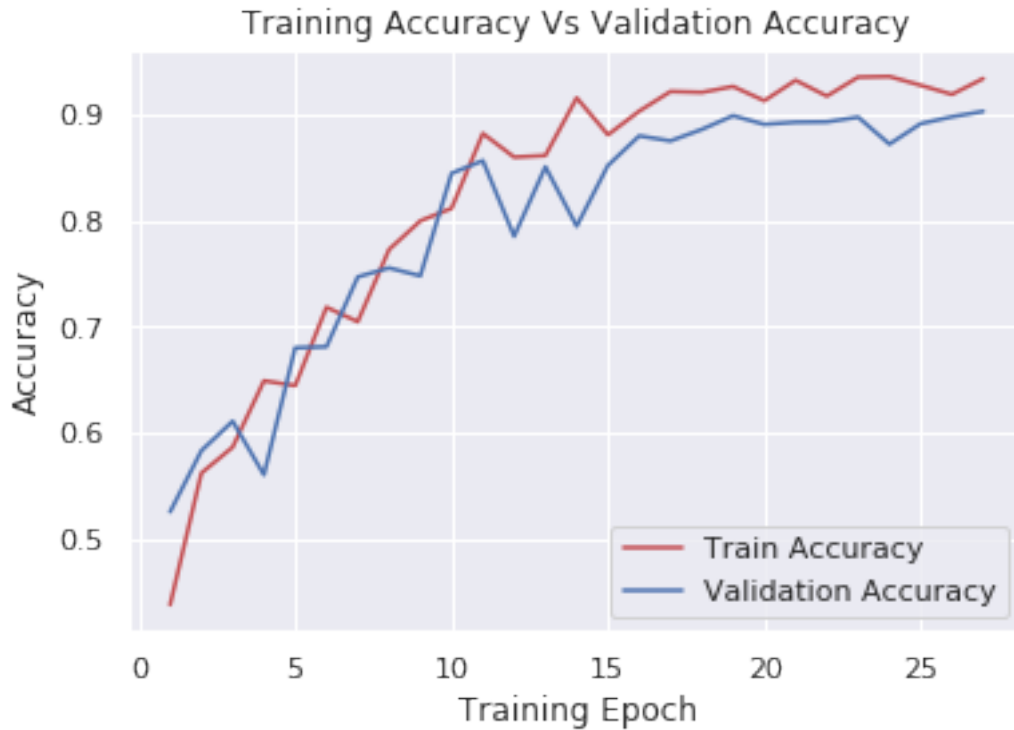
```
=====
Total params: 11,430
Trainable params: 11,430
Non-trainable params: 0
-----
```

None

```
In [28]: # Train the Model
        model_a2_name = 'Single_Layered_LSTM_2.h5'
        hist_model_a2 = train_model(model_a2, model_a2_name, X_train, y_train, num_epochs)

In [29]: # plot loss, accuracy curves
        plot_loss_curve(hist_model_a2)
        plot_accuracy_curve(hist_model_a2)
```





In [30]: # Evaluate the model on Test data

```
train_score_model_a2 = evaluate_model(model_a2, model_a2_name, X_train, y_train,
                                     title_suffix='Train', plot_fig=False)

train_score_model_a2 = [round(item, 4) for item in train_score_model_a2]
```

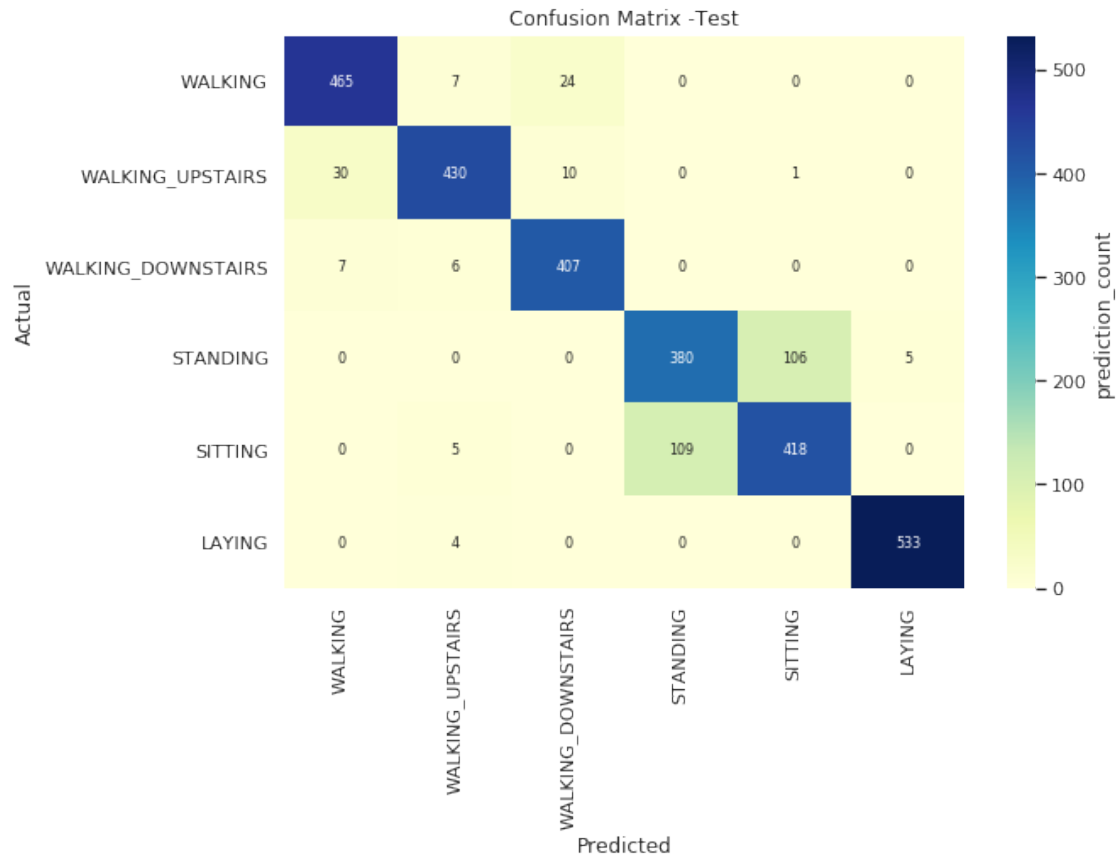
Restored best model weights from saved file ./model/Single_Layered_LSTM_2.h5
 7352/7352 [=====] - 2s 293us/step

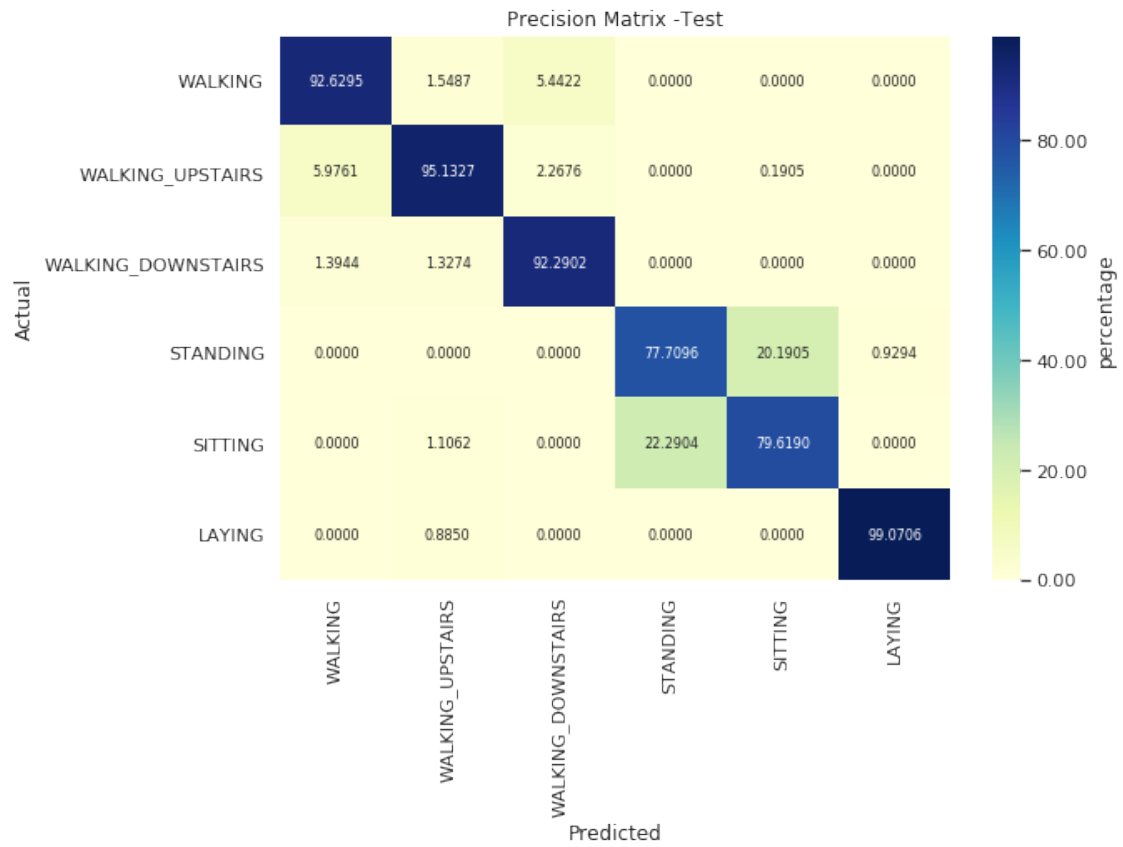
In [31]: # Evaluate the model on Test data

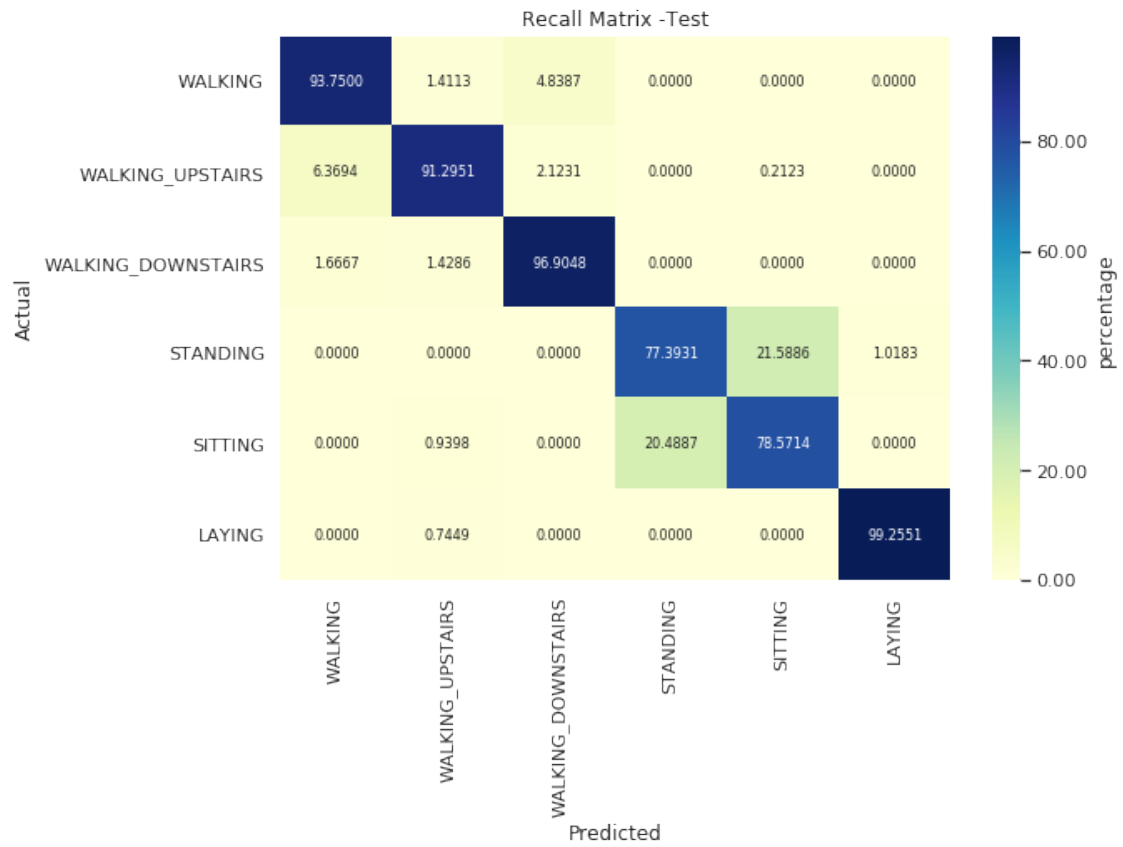
```
test_score_model_a2 = evaluate_model(model_a2, model_a2_name, X_test, y_test,
                                    title_suffix='Test', plot_fig=True)

test_score_model_a2 = [round(item, 4) for item in test_score_model_a2]
```

Restored best model weights from saved file ./model/Single_Layered_LSTM_2.h5
 2947/2947 [=====] - 1s 338us/step









```
In [32]: table_entry_model_a2 = ['1 Layered LSTM(2)', h_params_a2] + train_score_model_a2 + \
      test_score_model_a2
```

7.2 b) 2-Layered Architecture

7.2.1 Architecture 1

```
In [33]: h_params_b1 = (32, 18, 0.30, 0.45,)
      model_b1 = build_model_b(timesteps, input_dim, h_params_b1)
```

Layer (type)	Output Shape	Param #
lstm_3 (LSTM)	(None, 128, 32)	5376
dropout_3 (Dropout)	(None, 128, 32)	0
lstm_4 (LSTM)	(None, 18)	3672

dropout_4 (Dropout)	(None, 18)	0
dense_3 (Dense)	(None, 6)	114

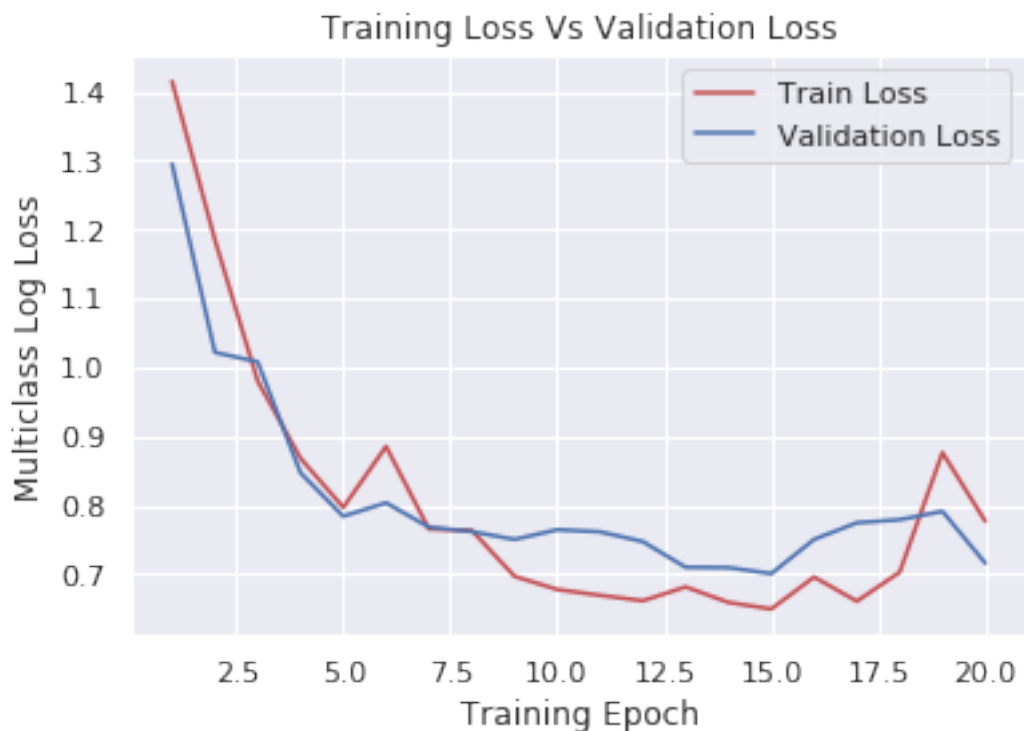
Total params: 9,162
 Trainable params: 9,162
 Non-trainable params: 0

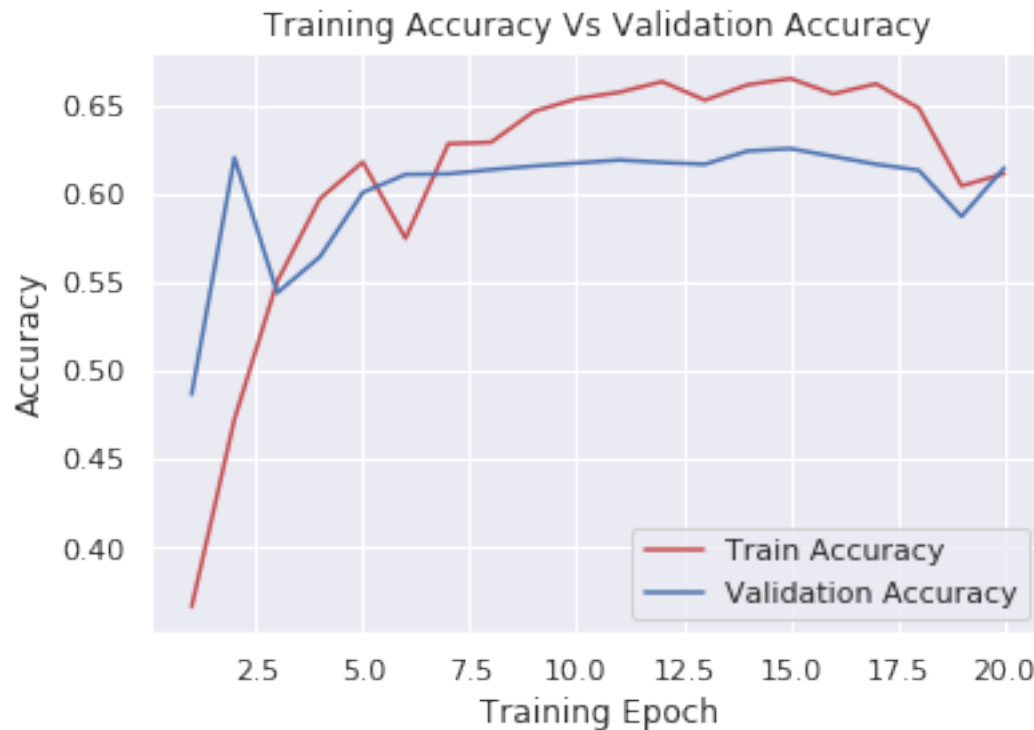
None

```

In [34]: # Train the Model
         model_b1_name = 'Two_Layered_LSTM_1.h5'
         hist_model_b1 = train_model(model_b1, model_b1_name, X_train, y_train, num_epochs)

In [35]: # plot loss, accuracy curves
         plot_loss_curve(hist_model_b1)
         plot_accuracy_curve(hist_model_b1)
  
```





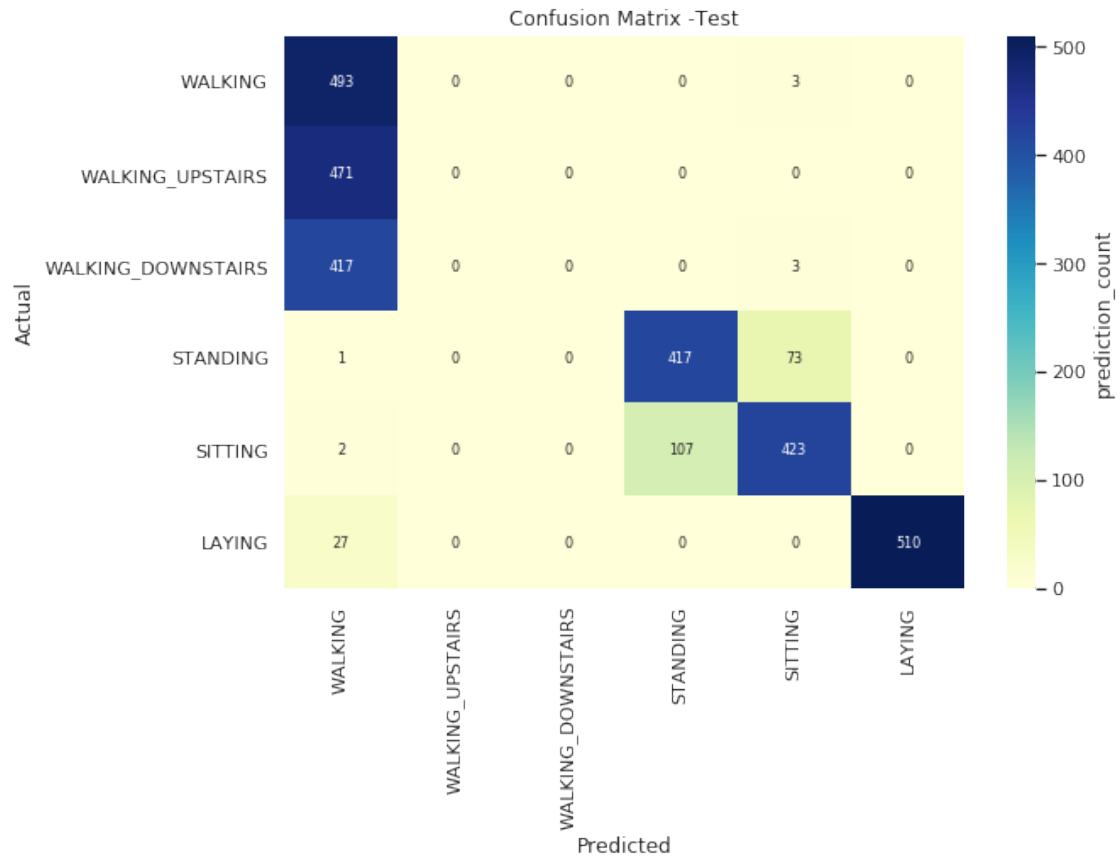
```
In [36]: # Evaluate the model on train data
train_score_model_b1 = evaluate_model(model_b1, model_b1_name, X_train, y_train,
                                     title_suffix='Train', plot_fig=False)
train_score_model_b1 = [round(item, 4) for item in train_score_model_b1]
```

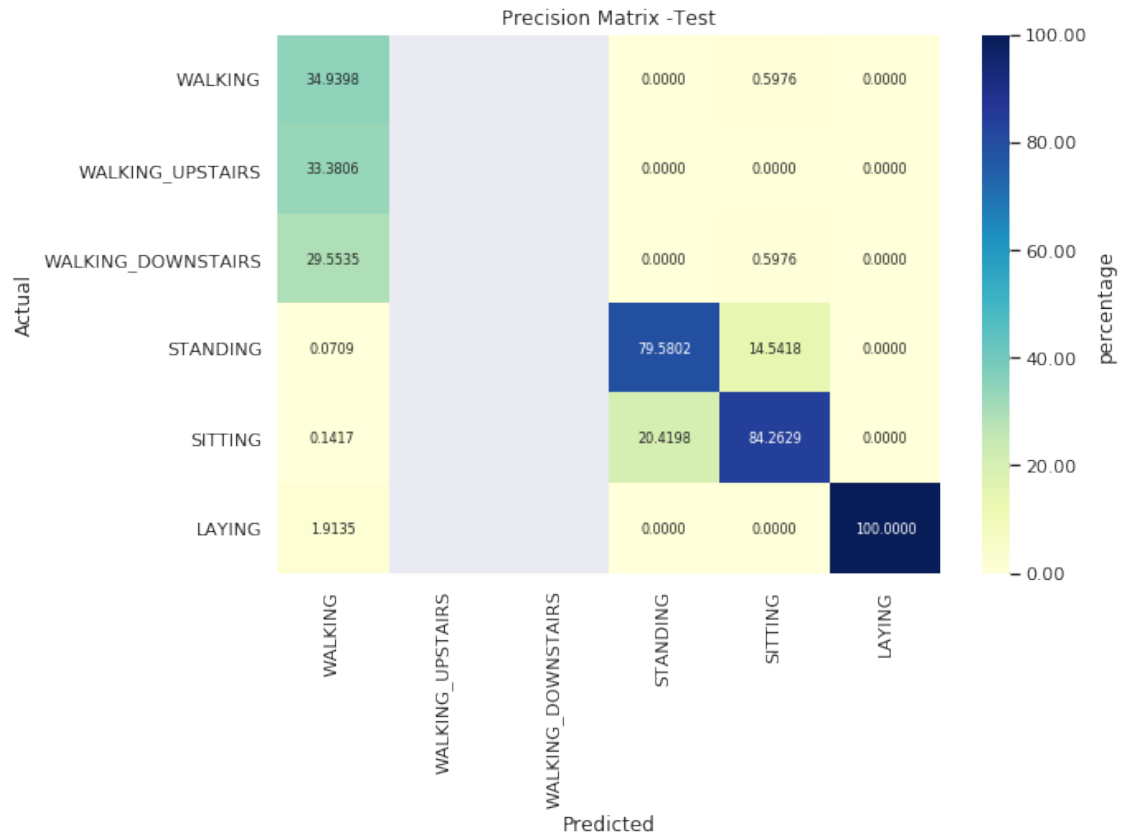
Restored best model weights from saved file ./model/Two_Layered_LSTM_1.h5
 7352/7352 [=====] - 4s 575us/step

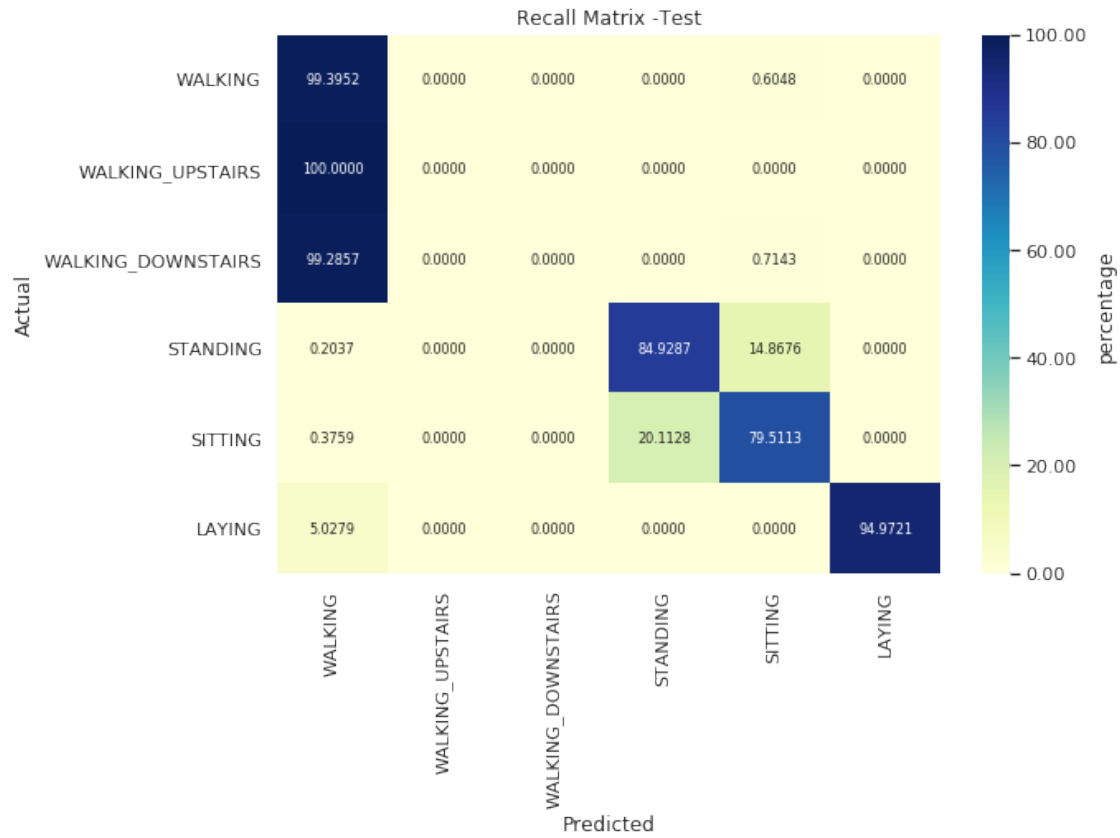
/home/nisheel-s/anaconda3/lib/python3.6/site-packages/sklearn/metrics/classification.py:1143: Un
 'precision', 'predicted', average, warn_for)

```
In [37]: # Evaluate the model on test data
test_score_model_b1 = evaluate_model(model_b1, model_b1_name, X_test, y_test,
                                    title_suffix='Test', plot_fig=True)
test_score_model_b1 = [round(item, 4) for item in test_score_model_b1]
```

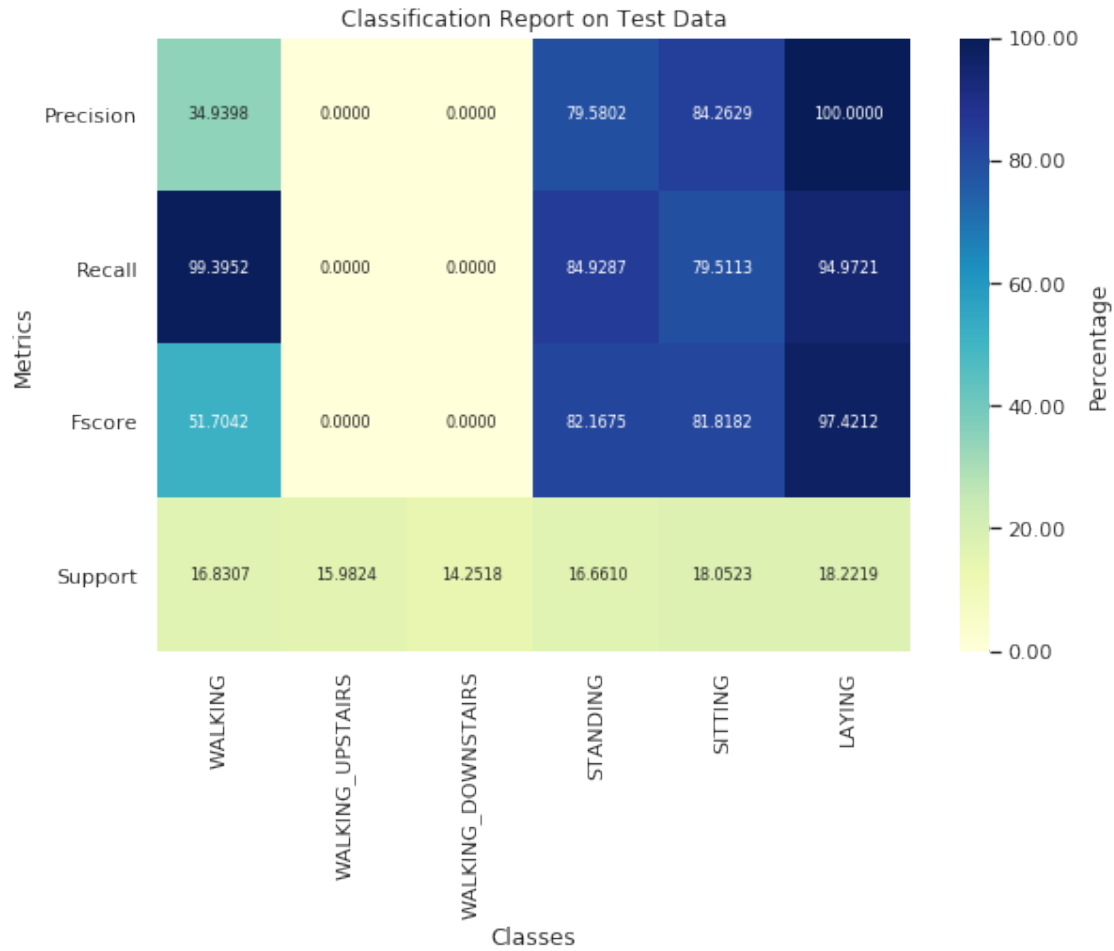
Restored best model weights from saved file ./model/Two_Layered_LSTM_1.h5
 2947/2947 [=====] - 2s 656us/step







```
/home/nisheel-s/anaconda3/lib/python3.6/site-packages/sklearn/metrics/classification.py:1143: Un
'precision', 'predicted', average, warn_for)
```



```
In [38]: table_entry_model_b1 = ['2 Layered LSTM(1)', h_params_b1] + train_score_model_b1 + \
      test_score_model_b1
```

7.2.2 Architecture 2

```
In [39]: h_params_b2 = (44, 23, 0.32, 0.48,)
      model_b2 = build_model_b(timesteps, input_dim, h_params_b2)
```

Layer (type)	Output Shape	Param #
lstm_5 (LSTM)	(None, 128, 44)	9504
dropout_5 (Dropout)	(None, 128, 44)	0
lstm_6 (LSTM)	(None, 23)	6256
dropout_6 (Dropout)	(None, 23)	0

```

-----
dense_4 (Dense)                (None, 6)                144
=====
Total params: 15,904
Trainable params: 15,904
Non-trainable params: 0
-----
None

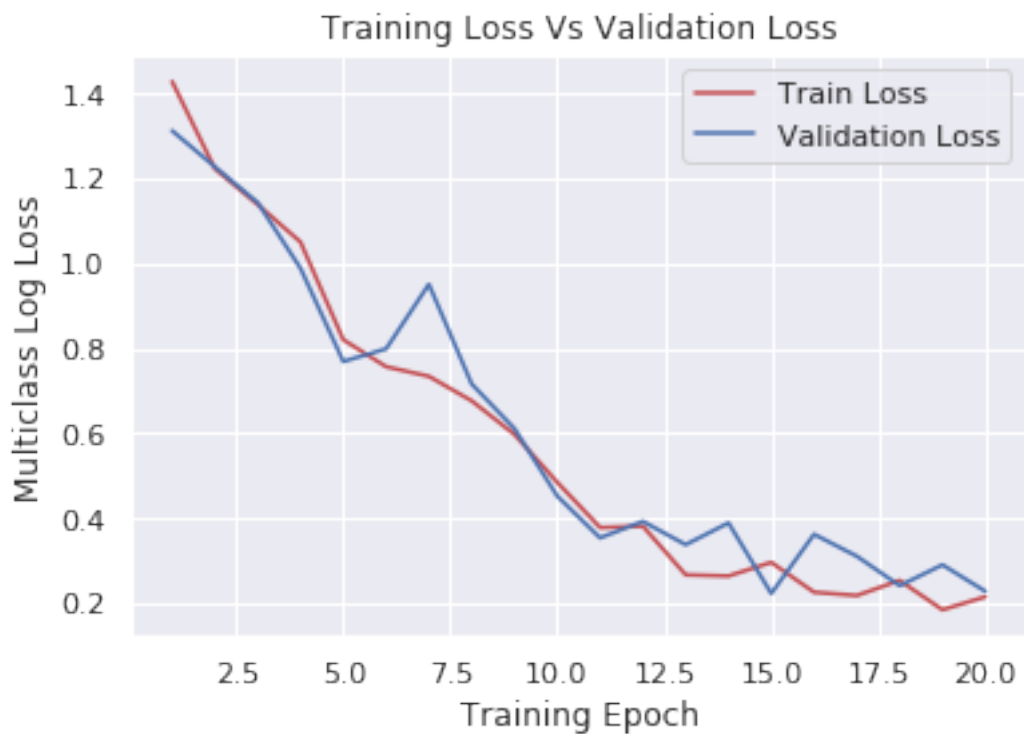
```

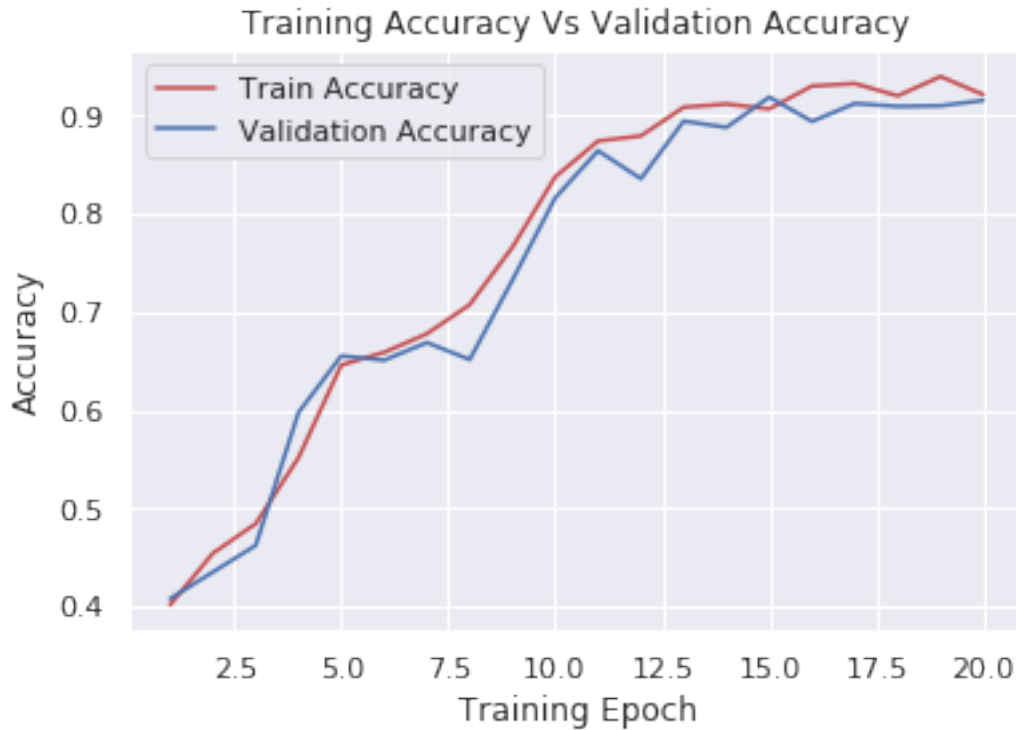
```

In [40]: # Train the Model
         model_b2_name = 'Two_Layered_LSTM_2.h5'
         hist_model_b2 = train_model(model_b2, model_b2_name, X_train, y_train, num_epochs)

In [41]: # plot loss, accuracy curves
         plot_loss_curve(hist_model_b2)
         plot_accuracy_curve(hist_model_b2)

```



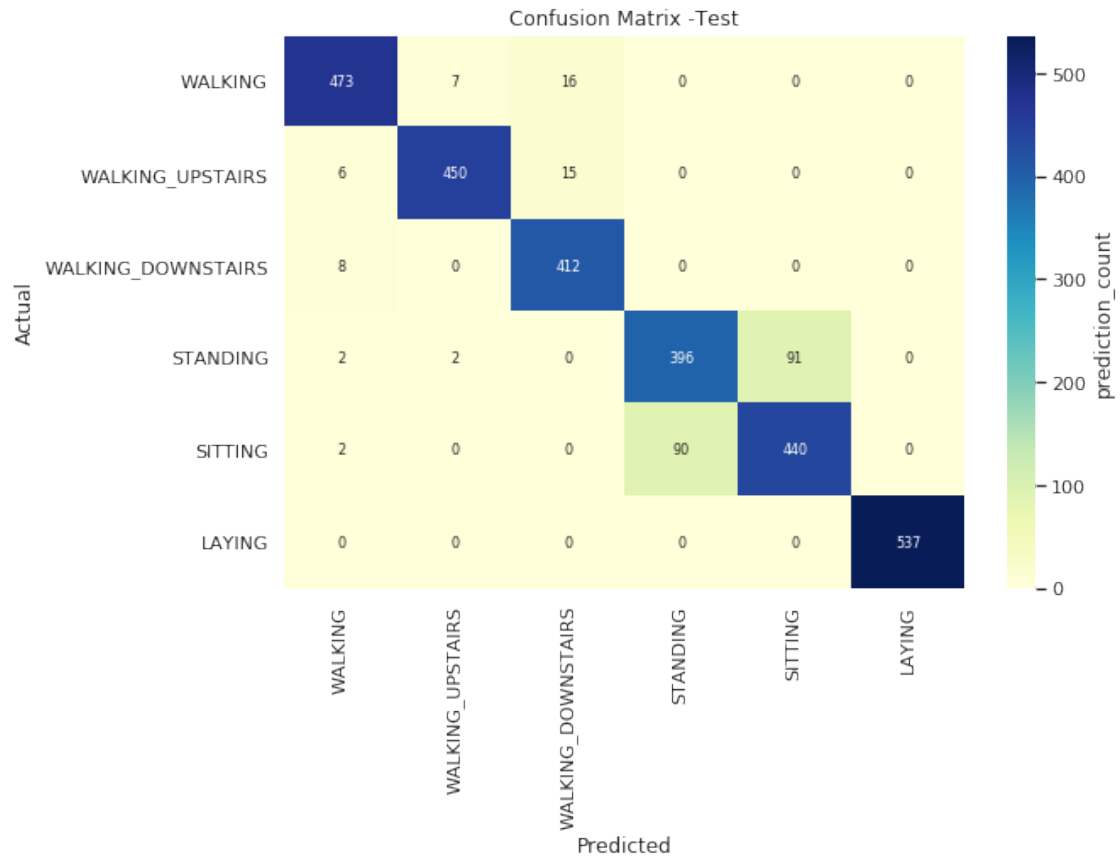


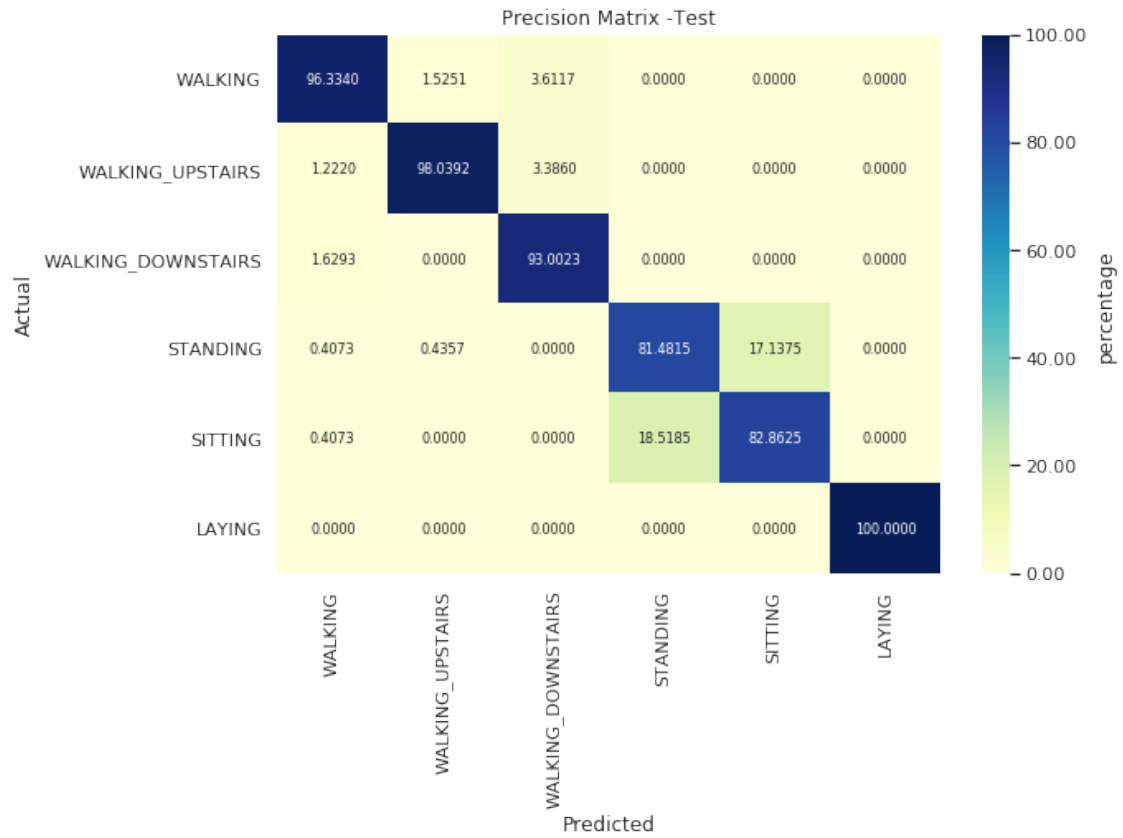
```
In [42]: # Evaluate the model on train data
train_score_model_b2 = evaluate_model(model_b2, model_b2_name, X_train, y_train,
                                     title_suffix='Train', plot_fig=False)
train_score_model_b2 = [round(item, 4) for item in train_score_model_b2]
```

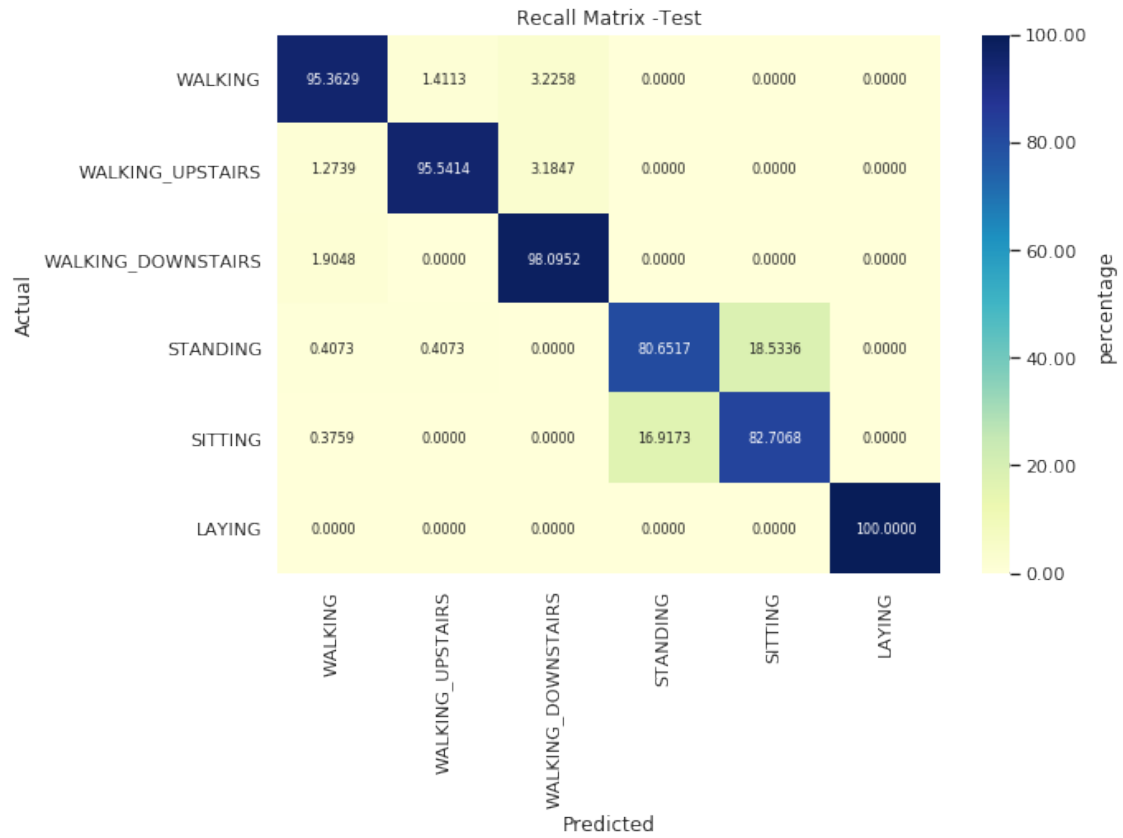
Restored best model weights from saved file ./model/Two_Layered_LSTM_2.h5
 7352/7352 [=====] - 5s 687us/step

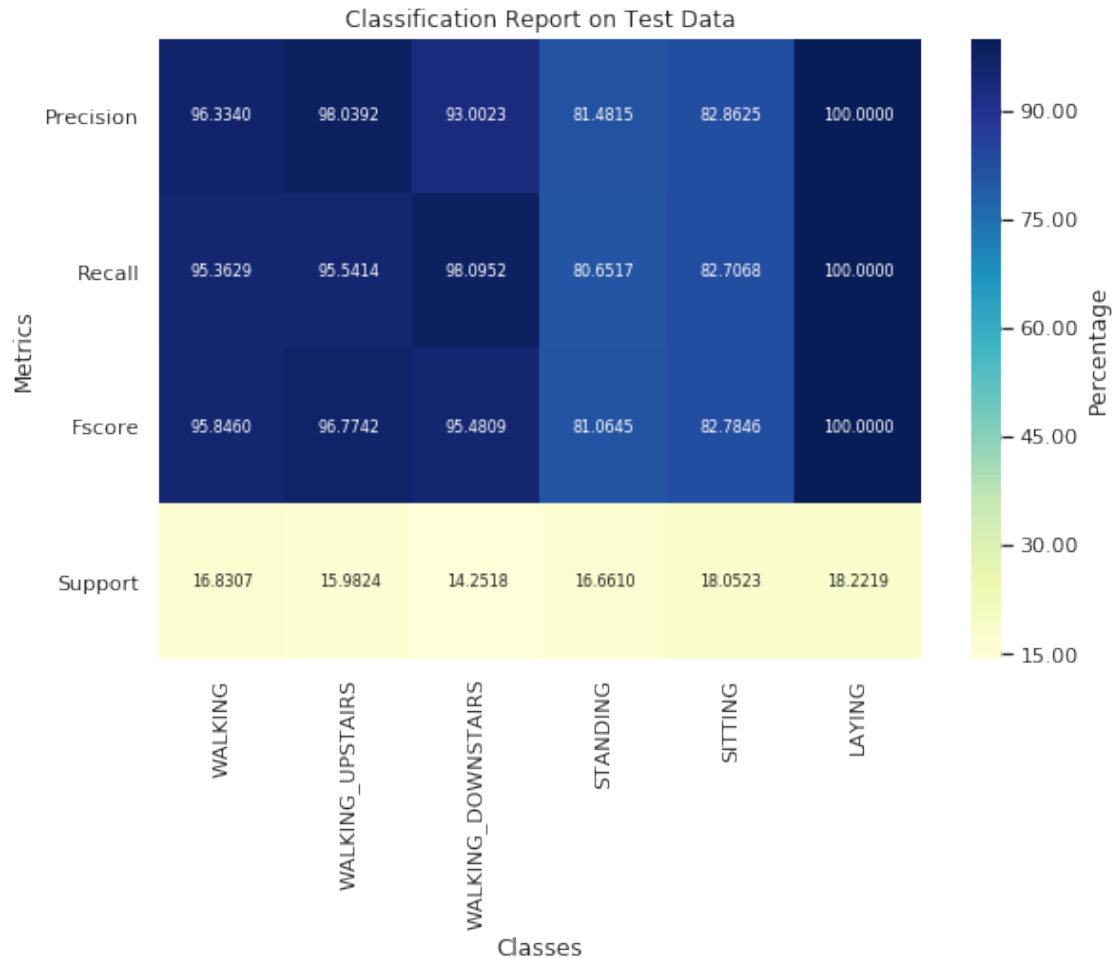
```
In [43]: # Evaluate the model on test data
test_score_model_b2 = evaluate_model(model_b2, model_b2_name, X_test, y_test,
                                    title_suffix='Test', plot_fig=True)
test_score_model_b2 = [round(item, 4) for item in test_score_model_b2]
```

Restored best model weights from saved file ./model/Two_Layered_LSTM_2.h5
 2947/2947 [=====] - 2s 786us/step









```
In [44]: table_entry_model_b2 = ['2-Layered LSTM(2)', h_params_b2] + train_score_model_b2 + \
    test_score_model_b2
```

8 Results

```
In [45]: Pret_table = PrettyTable()
    Pret_table.field_names = ['Model', '"Hyperparam (#units,dropout)"',
    'Train Loss', 'Train Acc (%)', 'Test Loss',
    'Test Acc (%)']
    Pret_table.title = 'HAR LSTM Model Performances'

    Pret_table.add_row(table_entry_model_a1)
    Pret_table.add_row(table_entry_model_a2)
    Pret_table.add_row(table_entry_model_b1)
    Pret_table.add_row(table_entry_model_b2)
    print(Pret_table)
```

Model	Hyperparam (#units,dropout)	Train Loss	Train Acc (%)	Test Loss	Test Acc (%)
1 Layered LSTM(1)	(32, 0.32)	0.1624	93.7432	0.3532	81.89
1 Layered LSTM(2)	(48, 0.4)	0.1592	94.056	0.3041	81.89
2 Layered LSTM(1)	(32, 18, 0.3, 0.45)	0.6048	67.519	0.7012	61.89
2-Layered LSTM(2)	(44, 23, 0.32, 0.48)	0.1456	94.9674	0.2236	91.89

9 Procedure Summary

The dataset is re-formatted to feed into LSTM model

The dataset is partitioned based on the number of users , 21 users in train side, 9 in test side (70:30) split

Two versions of single layered LSTM architecture is designed

Two versions of two layered LSTM architecture is designed

Train vs validation loss plot is made to evaluate performance

10 Conclusion

2-Layered LSTM architecture 2 performed well with 91.89 % accuracy

2-Layered LSTM architecture performance is not good as the accuracy is 62% very less compared to other models

Single layered architecture 1 showed 7% deviation between train, test. It shows tendency to overfit

Hyperparam of 2-layered architecture 2 can be tuned to improve the results further

All models get confused more in classifying between sitting & standing