Import required liberaries

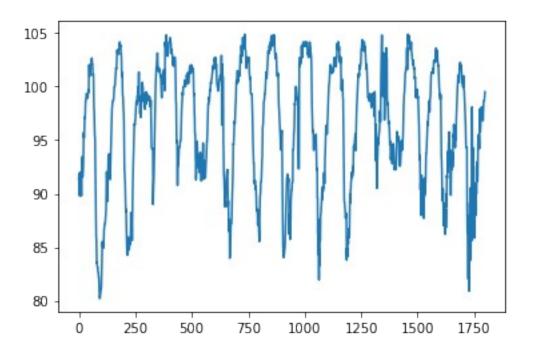
```
#import liberaries
import numpy as np
import pandas as pd
from pandas import read_csv
from numpy import nan
from numpy import isnan
from pandas import to_numeric
import matplotlib.pyplot as plt
```

load dataset

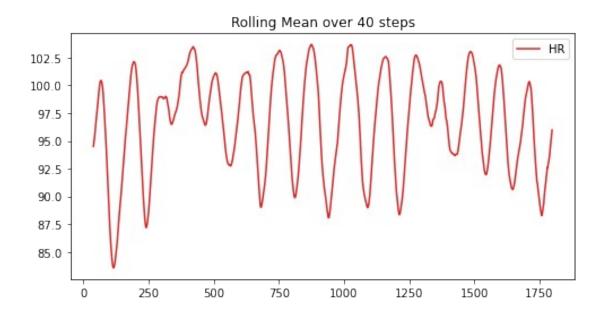
```
# load data
dataset = read_csv('heart-rate-time-series.csv')
dataset.columns=['HR'] #make heading for the column
dataset.head()
        HR
0 91.4634
1 91.1834
2 91.8788
3 91.1772
4 89.7992
from numpy import nan
from numpy import isnan
from pandas import to numeric
# mark all missing values
dataset.replace('?', nan, inplace=True)
# make dataset numeric
dataset = dataset.astype('float32')
dataset.head()
          HR
0 91.463402
1 91.183403
2 91.878799
3 91.177200
4 89.799202
```

Plot dataset to get more information about dataset plt.plot(dataset)

```
[<matplotlib.lines.Line2D at 0xd89007fc70>]
```



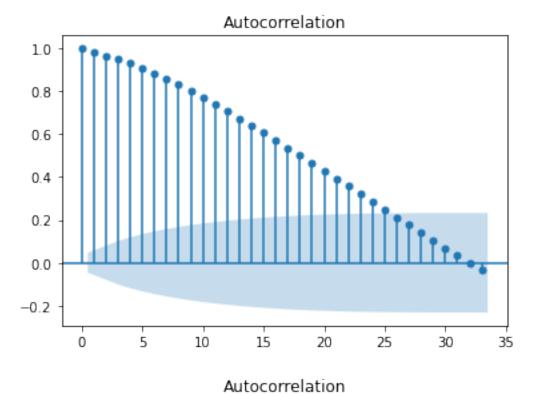
Moving averaging over the dataset to make it smoother dataset.rolling(window = 40).mean().plot(figsize=(8,4), color="tab:red", title="Rolling Mean over 40 steps");

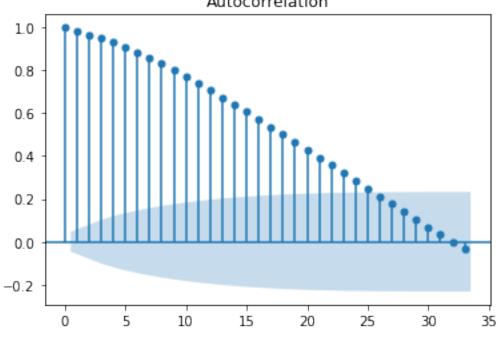


Do some statistics to got more insights

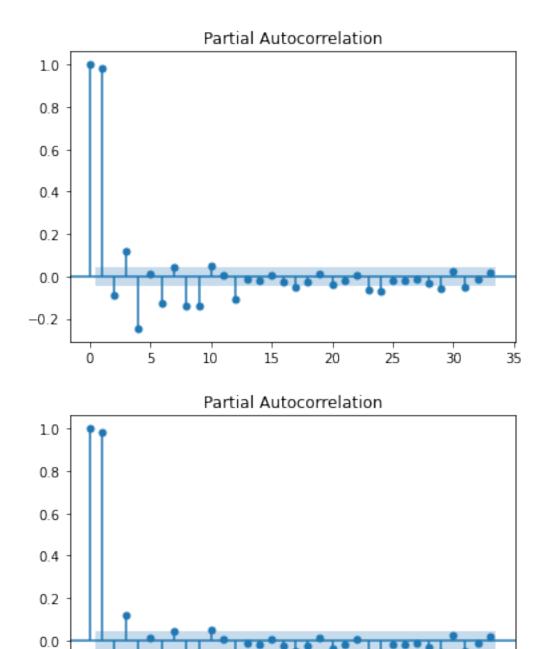
from statsmodels.graphics.tsaplots import plot_acf

plot_acf(dataset) #Auto Correlation Function ACF Plot or Autocorrelation plot





from statsmodels.graphics.tsaplots import plot_pacf
plot_pacf(dataset) #partial autocorrelation function (PACF) plot



Dicky Fuller test to see whether the time series is stationary or not from statsmodels.tsa.stattools import adfuller

```
dftest = adfuller(dataset['HR'], autolag = 'AIC')
print("1. ADF : ",dftest[0])
```

-0.2

Ó

```
print("2. P-Value : ", dftest[1])
print("3. Num Of Lags : ", dftest[2])
print("4. Num Of Observations Used For ADF Regression and Critical
Values Calculation :", dftest[3])
print("5. Critical Values :")
for key, val in dftest[4].items():
    print("\t",key, ": ", val)
1. ADF : -7.457988233959135
2. P-Value : 5.447866825089469e-11
3. Num Of Lags: 23
4. Num Of Observations Used For ADF Regression and Critical Values
Calculation: 1775
5. Critical Values :
      1%: -3.4340394547116797
      5%: -2.863169689048283
      10%: -2.5676375957944853
Differencing to make the time series stationary
# from pandas import Series
# # create a differenced series
# def difference(dataset, interval=10):
      diff = list()
      for i in range(interval, len(dataset)):
          value = dataset[i] - dataset[i - interval]
          diff.append(value)
      return Series(diff)
# dataset=difference(dataset, interval=10)
Normalizing the dataset
# Normalize time series data
from sklearn.preprocessing import MinMaxScaler
# prepare data for normalization
values=dataset.to numpy()
values = values.reshape((len(values), 1))
# train the normalization
scaler = MinMaxScaler(feature range=(-1, 1))
scaler = scaler.fit(values)
print('Min: %f, Max: %f' % (scaler.data_min_, scaler.data_max_))
# normalize the dataset
dataset = scaler.transform(values)
dataset
Min: 80.213898, Max: 104.894997
array([[-0.08841181],
```

[-0.11110115],

```
[-0.05475044],
       [ 0.52234745],
       [ 0.5488291 ],
       [ 0.5644846 ]], dtype=float32)
dataset.shape
(1799, 1)
Transform time series data to supervised data
from pandas import DataFrame
from pandas import concat
def series to supervised(data, n in=360, n out=600, dropnan=True):
    Frame a time series as a supervised learning dataset.
    Arguments:
        data: Sequence of observations as a list or NumPy array.
        n in: Number of lag observations as input (X).
        n out: Number of observations as output (y).
        dropnan: Boolean whether or not to drop rows with NaN values.
    Returns:
        Pandas DataFrame of series framed for supervised learning.
    n vars = 1 if type(data) is list else data.shape[1]
    df = DataFrame(data)
    cols, names = list(), list()
    # input sequence (t-n, \ldots t-1)
    for i in range(n in, 0, -1):
        cols.append(df.shift(i))
        names += [('var%d(t-%d)' % (j+1, i)) for j in range(n_vars)]
    # forecast sequence (t, t+1, \ldots t+n)
    for i in range(0, n out):
        cols.append(df.shift(-i))
        if i == 0:
            names += [('var%d(t)' % (j+1)) for j in range(n vars)]
        else:
            names += [('var%d(t+%d)' % (j+1, i)) for j in
range(n vars)]
    # put it all together
    agg = concat(cols, axis=1)
    agg.columns = names
    # drop rows with NaN values
    if dropnan:
        agg.dropna(inplace=True)
    return agg
```

data = series_to_supervised(dataset, 360, 600) #transform time series
dataset to supervised datasets
print(data)

,	var1(t-360)	var1(t-359)	var1(t-358)	var1(t-357)	var1(t-356)
\ 360	-0.088412	-0.111101	-0.054750	-0.111604	-0.223268
361	-0.111101	-0.054750	-0.111604	-0.223268	-0.178059
362	-0.054750	-0.111604	-0.223268	-0.178059	-0.043090
363	-0.111604	-0.223268	-0.178059	-0.043090	-0.052182
364	-0.223268	-0.178059	-0.043090	-0.052182	-0.114537
1195	0.740126	0.790689	0.837852	0.872210	0.873831
1196	0.790689	0.837852	0.872210	0.873831	0.849196
1197	0.837852	0.872210	0.873831	0.849196	0.858191
1198	0.872210	0.873831	0.849196	0.858191	0.882744
1199	0.873831	0.849196	0.858191	0.882744	0.906731
	var1(t-355) \	var1(t-354)	var1(t-353)	var1(t-352)	var1(t-351)
360	-0.178059	-0.043090	-0.052182	-0.114537	-0.048406
361	-0.043090	-0.052182	-0.114537	-0.048406	-0.055107
362	-0.052182	-0.114537	-0.048406	-0.055107	-0.193666
363	-0.114537	-0.048406	-0.055107	-0.193666	-0.222822
364	-0.048406	-0.055107	-0.193666	-0.222822	-0.073583
 1195	0.849196	0.858191	0.882744	0.906731	0.916292
1196	0.858191	0.882744	0.906731	0.916292	0.892306
1197 	0.882744	0.906731	0.916292	0.892306	0.893035

1198	0.906731	0.916292	0.892306	0.893035	0.922127
1199	0.916292	0.892306	0.893035	0.922127	0.938171
• • • •	- /	- /	- / >	- ()	- ()
\	var1(t+590)	var1(t+591)	var1(t+592)	var1(t+593)	var1(t+594)
360	0.127251	0.211470	0.248275	0.279165	0.291668
361	0.211470	0.248275	0.279165	0.291668	0.394192
362	0.248275	0.279165	0.291668	0.394192	0.476231
363	0.279165	0.291668	0.394192	0.476231	0.464732
364	0.291668	0.394192	0.476231	0.464732	0.503555
1195	0.418243	0.444036	0.416979	0.365726	0.344810
1196	0.444036	0.416979	0.365726	0.344810	0.396064
1197	0.416979	0.365726	0.344810	0.396064	0.466191
1198	0.365726	0.344810	0.396064	0.466191	0.492324
1199	0.344810	0.396064	0.466191	0.492324	0.496716
	var1(t+595)	var1(t+596)	var1(t+597)	var1(t+598)	var1(t+599)
360	0.394192	0.476231	0.464732	0.503555	0.568366
361	0.476231	0.464732	0.503555	0.568366	0.603340
362	0.464732	0.503555	0.568366	0.603340	0.603340
363	0.503555	0.568366	0.603340	0.603340	0.603340
364	0.568366	0.603340	0.603340	0.603340	0.588746
1195	0.396064	0.466191	0.492324	0.496716	0.496716
1196	0.466191	0.492324	0.496716	0.496716	0.496716

1197	0.492324	0.496716	0.496716	0.496716	0.522347
1198	0.496716	0.496716	0.496716	0.522347	0.548829
1199	0.496716	0.496716	0.522347	0.548829	0.564485

[840 rows \times 960 columns]

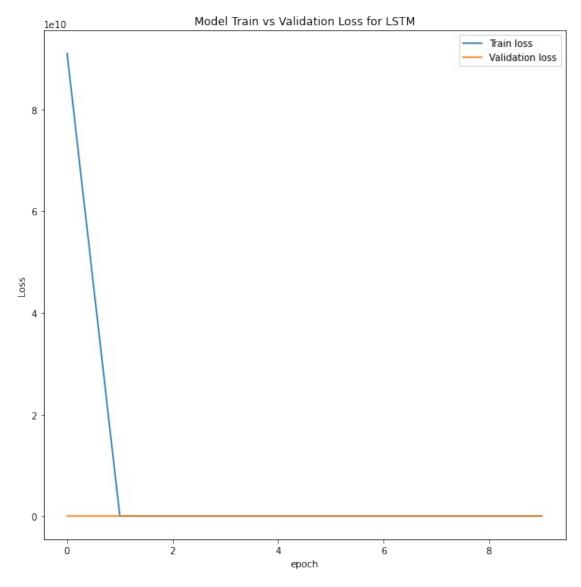
split dataset into train and test sets

```
# split into train and test sets
values = data.values
n train = int(len(values)*0.8) # 80% of data for training
train = values[:n train, :]
test = values[n train:, :]
n_train, train.shape, test.shape
(672, (672, 960), (168, 960))
np.info(train)
class: ndarray
shape: (672, 960)
strides: (4, 3360)
itemsize: 4
aligned: True
contiguous: False
fortran: False
data pointer: 0xd895a89040
byteorder: little
byteswap: False
type: float32
np.info(test)
class: ndarray
shape: (168, 960)
strides: (4, 3360)
itemsize: 4
aligned: True
contiguous: False
fortran: False
data pointer: 0xd895a89ac0
byteorder: little
byteswap: False
type: float32
np.info(train[:, -600:])
```

```
class: ndarray
shape: (672, 600)
strides: (4, 3360)
itemsize: 4
aligned: True
contiquous: False
fortran: False
data pointer: 0xd895bb0540
byteorder: little
byteswap: False
type: float32
split train and test sets into input and outputs
# split into input and outputs
train_X, train_y = train[:, :-600], train[:, -600:] #600 is the number
of outpout
test_X, test_y = test[:, :-600], test[:, -600:]
train X.shape, train y.shape
((672, 360), (672, 600))
prepare data to enter the network
# reshape input to be 3D [samples, timesteps, features]
features =1
train X = train X.reshape((train X.shape[0],
train X.shape[1],features))
test_X = test_X.reshape((test_X.shape[0], test_X.shape[1],features))
print(train X.shape, train y.shape, test X.shape, test y.shape)
(672, 360, 1) (672, 600) (168, 360, 1) (168, 600)
build the model
from keras.models import Sequential
from keras.layers import Dense
from keras.layers import Flatten
from keras.layers import LSTM
from keras.layers import Dropout
from keras import layers, callbacks
from keras.layers import Dense, LSTM, Dropout, GRU, Bidirectional
dropout=0.5
units=10
nout=600
# Create BiLSTM model
def create model bilstm(units):
```

```
model = Sequential()
    # First layer of BiLSTM
    model.add(Bidirectional(LSTM(units = units, activation='relu',
return sequences=True),input shape=(train X.shape[1],
train X.shape[2])))
    model.add(Dropout(dropout))
    # Second laver of BiLSTM
    #model.add(Bidirectional(LSTM(units = units)))
    model.add(Dense(nout))
    #Compile model
    model.compile(loss='mse', optimizer='adam')
    return model
# Create LSTM or GRU model
def create model(units, m):
    model = Sequential()
    # First laver of LSTM
    model.add(m (units = units, activation='relu', input shape =
[train X.shape[1], train X.shape[2]]))
    model.add(Dropout(dropout))
    # Second layer of LSTM
    #model.add(m (units = units))
    #model.add(Dropout(dropout))
    model.add(Dense(nout))
    #Compile model
    model.compile(loss='mse', optimizer='adam')
    return model
# BiLSTM
model bilstm = create model bilstm(units)
# GRU and LSTM
model gru = create model(units, GRU)
model lstm = create model(units, LSTM)
define a function to fit the model
#fit models
epoch, batch= 10,1
def fit model(model):
    history = model.fit(train X, train y, epochs=epoch,
batch size=batch, validation data=(test X, test y), verbose=1,
shuffle=False)
    return history
#history bilstm = fit model(model bilstm)
```

```
history lstm = fit model(model lstm)
#history gru = fit model(model gru)
Epoch 1/10
672/672 [============ ] - 41s 60ms/step - loss:
91032895488.0000 - val loss: 0.2115
Epoch 2/10
672/672 [============ ] - 40s 60ms/step - loss:
0.2500 - val loss: 0.2057
Epoch 3/10
672/672 [============ ] - 46s 68ms/step - loss:
0.2412 - val loss: 0.2029
Epoch 4/10
672/672 [============ ] - 49s 73ms/step - loss:
0.2398 - val loss: 0.2033
Epoch 5/10
672/672 [============ ] - 50s 75ms/step - loss:
0.2356 - val loss: 0.2037
Epoch 6/10
0.2353 - val loss: 0.2054
Epoch 7/10
672/672 [============ ] - 52s 77ms/step - loss:
0.2343 - val loss: 0.2066
Epoch 8/10
672/672 [============ ] - 53s 79ms/step - loss:
0.2294 - val loss: 0.2091
Epoch 9/10
672/672 [============ ] - 54s 80ms/step - loss:
0.2324 - val loss: 0.2093
Epoch 10/10
672/672 [============ ] - 53s 79ms/step - loss:
0.2303 - val_loss: 0.2098
plot the results
#plot train loss and validation loss
def plot_loss (history, model_name):
   plt.\overline{f}igure(figsize = (10, 10))
   plt.plot(history.history['loss'])
   plt.plot(history.history['val loss'])
   plt.title('Model Train vs Validation Loss for ' + model name)
   plt.ylabel('Loss')
   plt.xlabel('epoch')
   plt.legend(['Train loss', 'Validation loss'], loc='upper right')
#plot_loss (history_bilstm, 'BiLSTM')
plot_loss (history_lstm, 'LSTM')
#plot loss (history gru, 'GRU')
```



#model_bilstm.summary()
model_lstm.summary()
#model_gru.summary()

Model: "sequential_8"

Layer (type)	Output Shape	Param #
lstm_5 (LSTM)	(None, 10)	480
dropout_8 (Dropout)	(None, 10)	0
dense_8 (Dense)	(None, 600)	6600

Total params: 7,080 Trainable params: 7,080

```
make prediction
#make prediction
def prediction(model):
    prediction = model.predict(test X)
    return prediction
#prediction bilstm = prediction(model bilstm)
prediction lstm = prediction(model lstm)
prediction gru = prediction(model gru)
test X = test X.reshape((test X.shape[0], test X.shape[1]))
prediction lstm.shape, prediction gru.shape ,test X.shape
((168, 600), (168, 600), (168, 360))
rescale the predictions into actual scale and calculate RMSE
from numpy import concatenate
from math import sgrt
from sklearn.metrics import mean squared error
# invert scaling lstm
inv yhat lstm = concatenate((prediction lstm, test X[:, 1:]), axis=1)
inv yhat lstm = scaler.inverse transform(inv yhat lstm)
inv yhat lstm = inv yhat lstm[:,0]
# invert scaling gru
#inv yhat gru = concatenate((prediction gru, test X[:, 1:]), axis=1)
#inv yhat gru = scaler.inverse transform(inv yhat gru)
#inv yhat gru = inv yhat gru[:,0]
# invert scaling for actual
test_y = test_y.reshape((len(test_y), nout))
inv y = concatenate((test y, test X[:, 1:]), axis=1)
inv y = scaler.inverse transform(inv y)
inv y = inv y[:,0]
# calculate RMSE
rmse lstm = sqrt(mean squared error(inv y, inv yhat lstm))
#rmse gru = sqrt(mean squared error(inv y, inv yhat gru))
print('Test RMSE lstm: %.3f' % rmse lstm)
#print('Test RMSE gru: %.3f' % rmse gru)
inv yhat lstm.shape
Test RMSE lstm: 2.939
```

Define a function to calculate MAE and RMSE

```
# Define a function to calculate MAE and RMSE
def evaluate prediction(predictions, actual, model name):
    errors = predictions - actual
    mse = np.square(errors).mean()
    rmse = np.sqrt(mse)
    mae = np.abs(errors).mean()
    print(model name + ':')
    #print('Mean Absolute Error: {:.4f}'.format(mae))
    print('Root Mean Square Error: {:.4f}'.format(rmse))
    print('')
#evaluate prediction(prediction bilstm, test y, 'Bidirectional LSTM')
evaluate_prediction(prediction_lstm, test_y, 'LSTM')
#evaluate prediction(prediction gru, test y, 'GRU')
LSTM:
Root Mean Square Error: 0.4580
inv yhat lstm.shape, inv y.shape
((168,),(168,))
```

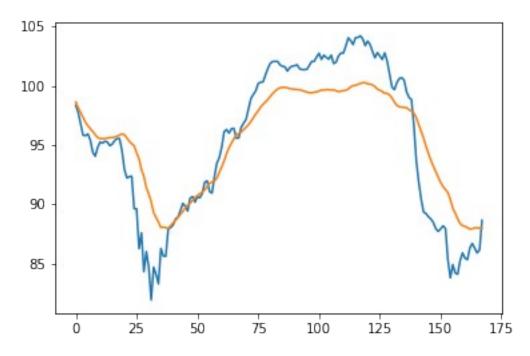
put actual values and predicted values alltogether in a dataframe and plot them

```
df = pd.DataFrame({'observation': inv_y, 'prediction':
inv_yhat_lstm} , columns=['observation', 'prediction']) #'gru
prediction2': inv_yhat_gru, 'bilstm prediction': inv_yhat_bilstm
df.head(20)
```

```
observation prediction
0
      98.344498
                 98.624763
1
      97.737602
                 98.139389
2
                 97.718750
      96.772797
3
      95.825600
                 97.314781
4
      95.791298
                 96.922714
5
      95.923599
                 96.610977
6
      95.394203
                 96.367989
7
     94.375702
                 96.128822
8
     94.054199
                 95.862381
9
     94.796303
                 95.629387
10
     95.244797
                 95.533890
```

```
11
      95.159203
                   95.535057
12
      95.311600
                   95.539604
13
      95.243896
                   95.583015
14
      94.941498
                   95.622971
15
      95.074600
                   95.627090
      95.376900
                   95.663689
16
17
      95.541397
                   95.746010
18
      95.541397
                   95.843689
19
      94.527000
                   95.930412
```

plt.plot(df)



Result Interpretion:

We can see from the above plot and dataframe that the predictions are very close to the actual values which is reasonable and shows that the network works correctly but it still could be more optimized