# **Test Your Algorithm**

# Instructions

1. From the Pulse Rate Algorithm Notebook you can do one of the following:

- Copy over all the **Code** section to the following Code block.
- Download as a Python ( . py ) and copy the code to the following Code block.
- 2. In the bottom right, click the Test Run button.

## **Didn't Pass**

If your code didn't pass the test, go back to the previous Concept or to your local setup and continue iterating on your algorithm and try to bring your training error down before testing again.

### Pass

If your code passes the test, complete the following! You **must** include a screenshot of your code and the Test being **Passed**. Here is what the starter filler code looks like when the test is run and should be similar. A passed test will include in the notebook a green outline plus a box with **Test passed**: and in the Results bar at the bottom the progress bar will be at 100% plus a checkmark with **All cells passed**.

#### Example

- Take a screenshot of your code passing the test, make sure it is in the format .png . If not a .png image, you will have to edit the Markdown render the image after Step 3. Here is an example of what the passed.png would look like
- 2. Upload the screenshot to the same folder or directory as this jupyter notebook.
- 3. Rename the screenshot to passed.png and it should show up below.



Passed

4. Download this jupyter notebook as a .pdf file.

5. Continue to Part 2 of the Project.

In [2]: import glob

```
from tgdm import tgdm
import numpy as np
import scipy as sp
import scipy.io
import scipy.signal
import os.path
from tqdm import tqdm
from sklearn.model selection import train test split
from sklearn.linear model import LinearRegression
from sklearn.model selection import KFold
from sklearn.ensemble import RandomForestRegressor,AdaBoostRegressor
from sklearn.metrics import mean squared error
def LoadTroikaDataset():
    0.0.0
    Retrieve the .mat filenames for the troika dataset.
    Review the README in ./datasets/troika/ to understand the organizat
ion of the .mat files.
    Returns:
        data fls: Names of the .mat files that contain signal data
        ref fls: Names of the .mat files that contain reference data
        <data fls> and <ref fls> are ordered correspondingly, so that r
ef fls[5] is the
            reference data for data fls[5], etc...
    ......
    data dir = "./datasets/troika/training data"
    data fls = sorted(glob.glob(data dir + "/DATA *.mat"))
    ref fls = sorted(glob.glob(data dir + "/REF *.mat"))
    return data fls, ref fls
def LoadTroikaDataFile(data fl):
```



```
Loads and extracts signals from a troika data file.
    Usage:
        data fls, ref fls = LoadTroikaDataset()
       ppg, accx, accy, accz = LoadTroikaDataFile(data fls[0])
   Args:
       data_fl: (str) filepath to a troika .mat file.
    Returns:
        numpy arrays for ppg, accx, accy, accz signals.
    0.0.0
   data = sp.io.loadmat(data fl)['sig']
    return data[2:]
def AggregateErrorMetric(pr_errors, confidence_est):
   Computes an aggregate error metric based on confidence estimates.
   Computes the MAE at 90% availability.
   Args:
       pr errors: a numpy array of errors between pulse rate estimates
and corresponding
            reference heart rates.
        confidence est: a numpy array of confidence estimates for each
pulse rate
            error.
    Returns:
        the MAE at 90% availability
    0.0.0
   # Higher confidence means a better estimate. The best 90% of the es
timates
    #
        are above the 10th percentile confidence.
    percentile90 confidence = np.percentile(confidence est, 10)
```



```
# Find the errors of the best pulse rate estimates
   best_estimates = pr_errors[confidence_est >= percentile90 confidenc
e1
    # Return the mean absolute error
    return np.mean(np.abs(best estimates))
def Evaluate():
   Top-level function evaluation function.
   Runs the pulse rate algorithm on the Troika dataset and returns an
aggregate error metric.
    Returns:
        Pulse rate error on the Troika dataset. See AggregateErrorMetri
с.
    0.0.0
    # Retrieve dataset files
   data fls, ref fls = LoadTroikaDataset()
   errs, conFs = [], []
   for data fl, ref fl in zip(data fls, ref fls):
       # Run the pulse rate algorithm on each trial in the dataset
        errors, confidence = RunPulseRateAlgorithm(data fl, ref fl)
       errs.append(errors)
        conFs.append(confidence)
       # Compute aggregate error metric
   errs = np.hstack(errs)
   conFs = np.hstack(conFs)
    return AggregateErrorMetric(errs, conFs)
def RunPulseRateAlgorithm(data fl, ref fl):
   Fs = 125 # Sample Frequency
   window len = 8 # Window to calculate PR
   window shift = 2 # Difference between windows
    reg, scores = Regressor()
   targets, features, sigs, subs = Data window8(data fl, ref fl)
```



```
error, confidence = [], []
    for i,feature in enumerate(features):
        est = reg.predict(np.reshape(feature, (1, -1)))[0]
        ppg, accx, accy, accz = sigs[i]
        ppg = Filter(ppg)
        accx = Filter(accx)
        accy = Filter(accy)
        accz = Filter(accz)
        n = len(ppq) * 3
        freq = np.fft.rfftfreq(n, 1/Fs)
        fft = np.abs(np.fft.rfft(ppg,n))
        fft[freq <= 40/60.0] = 0.0
        fft[freq \ge 240/60.0] = 0.0
        est Fs = est / 55.0
        Fs win = 30 / 60.0
        Fs win e = (freq >= est Fs - Fs win) & (freq <= est Fs +Fs win)
        conf = np.sum(fft[Fs win e])/np.sum(fft)
        error.append(np.abs((est-targets[i])))
        confidence.append(conf)
    return np.array(error), np.array(confidence)
def Data window8(data fl, ref fl):
    Fs=125 # Sampling frequency
    window len = 6 # Window to calculate PR
    window shift = 2 # Difference between windows
    sig = LoadTroikaDataFile(data fl)
    ref = scipy.io.loadmat(ref fl)["BPM0"]
    ref = np.array([x[0] for x in ref])
    subject name = os.path.basename(data fl).split('.')[0]
    start indxs, end indxs = Indexator(sig.shape[1], len(ref), Fs, wind
ow len, window shift)
    targets, features, sigs, subs = [], [], [], []
```



```
for i, s in enumerate(start indxs):
        start i = start indxs[i]
        end i = end indxs[i]
        ppg = sig[0, start i:end i]
        accx = sig[1, start i:end i]
        accy = sig[2, start i:end i]
        accz = sig[3, start i:end i]
        ppq = Filter(ppq)
        accx = Filter(accx)
        accy = Filter(accy)
        accz = Filter(accz)
        feature, ppg, accx, accy, accz = CreateFeature(ppg, accx, accy,
 accz)
        sigs.append([ppg, accx, accy, accz])
        targets.append(ref[i])
        features.append(feature)
        subs.append(subject name)
    return (np.array(targets), np.array(features), sigs, subs)
def Data window6():
    Fs=125 # Sampling rate
    window len = 6 # Window to calculate PR
    window shift = 2 # Difference between windows
    data fls, ref fls = LoadTroikaDataset()
    pbar = tqdm(list(zip(data fls, ref fls)), desc="Prepare Data")
    targets, features, sigs, subs = [], [], [], []
    for data fl, ref fl in pbar:
        sig = LoadTroikaDataFile(data fl)
        ref = scipy.io.loadmat(ref fl)["BPM0"]
        ref = np.array([x[0] for x in ref])
        subject name = os.path.basename(data fl).split('.')[0]
        start indxs, end indxs = Indexator(sig.shape[1], len(ref), Fs,
```



```
window_len,window_shift)
        for i, s in enumerate(start indxs):
            start i = start indxs[i]
            end i = end indxs[i]
            ppg = sig[0, start i:end i]
            accx = sig[1, start i:end i]
            accy = sig[2, start i:end i]
            accz = sig[3, start i:end i]
            ppq = Filter(ppq)
            accx = Filter(accx)
            accy = Filter(accy)
            accz = Filter(accz)
            feature, ppg, accx, accy, accz = CreateFeature(<math>ppg, accx, a
ccy, accz)
            sigs.append([ppg, accx, accy, accz])
            targets.append(ref[i])
            features.append(feature)
            subs.append(subject name)
    return (np.array(targets), np.array(features), sigs, subs)
def CreateFeature(ppg, accx, accy, accz):
    """ Create features """
    ppg = Filter(ppg)
    accx = Filter(accx)
   accy = Filter(accy)
    accz = Filter(accz)
    Fs = 125
   n = len(ppq) * 4
   freq = np.fft.rfftfreq(n, 1/Fs)
    fft = np.abs(np.fft.rfft(ppg,n))
    fft[freq <= 40/60.0] = 0.0
```



```
fft[freg \ge 240/60.0] = 0.0
    acct = np.sqrt(accx**2 + accy**2 + accz**2) # Total signal of acc
    acc fft = np.abs(np.fft.rfft(acct, n))
    acc fft[freq <= 40/60.0] = 0.0
    acc fft[freq >= 240/60.0] = 0.0
    ppg feature = freg[np.argmax(fft)]
    acc feature = freg[np.argmax(acc fft)]
    return (np.array([ppg feature, acc feature]), ppg, accx, accy, accz
def RegressionAlg(features, targets, subs):
    """ The rearession model"""
    AdaBoostRegressor
    regression = RandomForestRegressor(n estimators=400, max depth=16)
    scores = []
   lf = KFold(n splits=5)
    splits = lf.split(features,targets,subs)
    for i, (train idx, test idx) in enumerate(splits):
        X train, y train = features[train idx], targets[train idx]
        X test, y test = features[test_idx], targets[test_idx]
        regression.fit(X train, y train)
        y pred = regression.predict(X test)
        score = Error(y test, y pred)
        scores.append(score)
    return (regression, scores)
def Filter(signal):
    """Bandpass filter between 40 and 240 BPM"""
    pass band=(40/60.0, 240/60.0)
    Fs = 125
    b, a = scipy.signal.butter(3, pass band, btype='bandpass', fs=Fs)
    return scipy.signal.filtfilt(b, a, signal)
def Indexator(sig len, ref len, Fs=125, window len s=10, window shift s
```



```
=2):
    0.0.0
    Find start and end index to iterate over a set of signals
    ......
    # Set the length of the biggest signal with regards to the referenc
e signal
    if ref len < sig len:</pre>
        n = ref len
    else:
        n = sig len
    # Start Indexes
    start indxs = (np.cumsum(np.ones(n) * Fs * window shift s) - Fs * w
indow shift s).astype(int)
    # End Indexes (same size as the start indexes array)
    end indxs = start indxs + window len s * Fs
    return (start indxs, end indxs)
def Predict(reg,feature, ppg, accx, accy, accz):
    """Predict based on the regressor"""
    est = reg.predict(np.reshape(feature, (1, -1)))[0]
def Error(y test, y pred):
    .. .. ..
    Calculate error score of a Prediction
    return mean squared error(y test, y pred)
def Regressor():
    fname = "outfile.npy"
    req, scores = [], []
    if os.path.isfile(fname):
        [reg,scores] = np.load(fname,allow pickle=True)
        #
    else:
        targets, features, sigs, subs = Data window6()
        reg, scores = RegressionAlg(features, targets, subs)
```



	<pre>np.save("outfile", [reg,scores]) return reg, scores</pre>
In [ ]:	

