

DSRM

January 6, 2022

1 Introduction

The dataset used is PAMAP2 which is an Activity Monitoring dataset that covers 18 different physical activities which are taken by 9 different subjects, 8 men and 1 woman, taken using 3 inertial measurement units and a heart rate monitor. The dataset can be used for activity recognition and intensity estimation, while developing and applying algorithms of data processing, extraction and classification. the PAMAP2 Physical Activity Monitoring dataset will be used to extract actionable insights that will allow the development of HAR software and/or hardware for fitness tracking.

The objectives of the assignment are the following: - To carry out thorough exploratory data analysis and appropriately handle missing or dirty data; - To develop and test at least one hypothesis for a relationship between a single pair of attributes; - To develop and test at least one model which uses multiple attributes to make predictions.

2 Data Loading

The first step is to import all the necessary libraries

```
[2]: #Required cell: This cell needs to be executed to import the necessary libraries
import pandas as pd
from matplotlib import pyplot as plt
from matplotlib.colors import LinearSegmentedColormap
import seaborn as sns
import numpy as np
from scipy import stats
from scipy import integrate
from IPython.display import HTML, display
from scipy.stats import norm
from scipy.stats import t as the
from sklearn import svm
from sklearn.metrics import classification_report, accuracy_score, \
    precision_score, recall_score, f1_score
from sklearn import tree
%matplotlib inline
pd.set_option('display.max_rows', 20)
pd.set_option('display.max_columns', 70)
from sklearn.model_selection import train_test_split
```

As part of loading we have to create a list first (list_of_files) which consists of all the file names and the path which needs to be loaded. This list will be used for calling the files. Then we create a list, subjectID which consist of all numbers 1 to 9 which represents the 9 subjects. After that we create a dictionary which consists of the activity ID as the key and the name of the activity as the value. Then we create a list colNames which consists of the first 3 columns, then the IMU hand, chest ankle lists are created which consists of all the attributes which each of the IMU have. After create these lists, we concatnate them to the variable columns.

```
[3]: # Load data
list_of_files = ['D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/
↳subject101.dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject102.
↳dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject103.
↳dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject104.
↳dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject105.
↳dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject106.
↳dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject107.
↳dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject108.
↳dat',
    'D:/DS/DSRM/ASSIGNMENT/PAMAP2_Dataset/PAMAP2_Dataset/Protocol/subject109.
↳dat' ]

subjectID = [1,2,3,4,5,6,7,8,9]

activityIDdict = {0: 'transient',
    1: 'lying',
    2: 'sitting',
    3: 'standing',
    4: 'walking',
    5: 'running',
    6: 'cycling',
    7: 'Nordic_walking',
    9: 'watching_TV',
    10: 'computer_work',
    11: 'car driving',
    12: 'ascending_stairs',
    13: 'descending_stairs',
    16: 'vacuum_cleaning',
    17: 'ironing',
    18: 'folding_laundry',
    19: 'house_cleaning',
```

```

        20: 'playing_soccer',
        24: 'rope_jumping' }

colNames = ["timestamp", "activityID", "heartrate"]

IMUhand = ['handTemperature',
            'handAcc16_1', 'handAcc16_2', 'handAcc16_3',
            'handAcc6_1', 'handAcc6_2', 'handAcc6_3',
            'handGyro1', 'handGyro2', 'handGyro3',
            'handMagne1', 'handMagne2', 'handMagne3',
            'handOrientation1', 'handOrientation2', 'handOrientation3',
            ↪ 'handOrientation4']

IMUchest = ['chestTemperature',
            'chestAcc16_1', 'chestAcc16_2', 'chestAcc16_3',
            'chestAcc6_1', 'chestAcc6_2', 'chestAcc6_3',
            'chestGyro1', 'chestGyro2', 'chestGyro3',
            'chestMagne1', 'chestMagne2', 'chestMagne3',
            'chestOrientation1', 'chestOrientation2', 'chestOrientation3',
            ↪ 'chestOrientation4']

IMUankle = ['ankleTemperature',
            'ankleAcc16_1', 'ankleAcc16_2', 'ankleAcc16_3',
            'ankleAcc6_1', 'ankleAcc6_2', 'ankleAcc6_3',
            'ankleGyro1', 'ankleGyro2', 'ankleGyro3',
            'ankleMagne1', 'ankleMagne2', 'ankleMagne3',
            'ankleOrientation1', 'ankleOrientation2', 'ankleOrientation3',
            ↪ 'ankleOrientation4']

columns = colNames + IMUhand + IMUchest + IMUankle #all columns in one list

```

The function `load_activity_map()` is created which will be used to map the Id with the activity, the dunction `pd_fast_plot` is created which will be used to plot the graphs, when we create three lists, `light_acts`, `mod_acts`, `vig_acts` which has th eIDs of the activities which are light, moderate and vigorous and these are put correspondingly to the activily levels.

```

[4]: def load_activity_map():
        map = {}
        map[0] = 'transient'
        map[1] = 'lying'
        map[2] = 'sitting'
        map[3] = 'standing'
        map[4] = 'walking'
        map[5] = 'running'

```

```

map[6] = 'cycling'
map[7] = 'Nordic_walking'
map[9] = 'watching_TV'
map[10] = 'computer_work'
map[11] = 'car driving'
map[12] = 'ascending_stairs'
map[13] = 'descending_stairs'
map[16] = 'vacuum_cleaning'
map[17] = 'ironing'
map[18] = 'folding_laundry'
map[19] = 'house_cleaning'
map[20] = 'playing_soccer'
map[24] = 'rope_jumping'
return map
def pd_fast_plot(pd,column_a,column_b,title, figsize=(10,6)):
    plt.rcParams.update({'font.size': 16})
    size = range(len(pd))
    f, ax = plt.subplots(figsize=figsize)
    plt.bar(size, pd[column_a], color=plt.cm.Paired(size))
    a = ax.set_xticklabels(pd[column_b])
    b = ax.legend(fontsize = 20)
    c = ax.set_xticks(np.arange(len(pd)))
    d = ax.set_title(title)
    plt.show()

    #lying, sitting, standing and ironing
    light_acts = [1,2,3,17]
    #vacuum cleaning, descending stairs, walking, Nordic walking and cycling
    mod_acts = [16,13,4,7,6]
    #ascending stairs, running and rope jumping
    vig_acts = [12,5,24]

#Function used to classify activities
def map_met(act_id):
    if act_id in light_acts:
        return 'light'
    if act_id in mod_acts:
        return 'moderate'
    if act_id in vig_acts:
        return 'vigorous'

```

In the following cell we are loading all the data into a single dataframe, we call the files one by one and then put the content into a dataframe where we set the column names too.

```

[5]: dataCollection = pd.DataFrame()
    for file in list_of_files:

```

```

procData = pd.read_table(file, header=None, sep='\s+')
procData.columns = columns
procData['subject_id'] = int(file[-5])
dataCollection = dataCollection.append(procData, ignore_index=True)

dataCollection.reset_index(drop=True, inplace=True)

```

```
[6]: len(dataCollection)
```

```
[6]: 2872533
```

2.1 Data Cleaning

As part of cleaning we are dropping the unnecessary columns that were mention in the report, all the orientation attributes that is the hand, chest and the ankle one are removed. Then we remove the throo hand acceleration columns wfor 6g cause due to high impacts caused by certain movements (e.g. during running) with acceleration over 6g, it gets saturated sometimes. Hence the acceleration over 16g is recommended. Then we drop all the rows where the activity id is 0 since this data mainly covers transient activities between performing different activities, e.g. going from one location to the next activity's location, or waiting for the preparation of some equipment.

Then we create a new column activity level where we mention wether its a light, moderate or vigorous activity. After that we will create new rows where we combine the points in 3d space, we will make a combined column for handacc16_1,handacc16_2,handacc16_3 and likewise for handgyro and handmag. After this we will do the samefor ankle and chest. These will be helpfull while plotting.

```

[7]: #Data cleaning
dataCollection=dataCollection.
    ↳drop(['handAcc6_1','handAcc6_2','handAcc6_3','handOrientation1',
    ↳'handOrientation2', 'handOrientation3',
    ↳'handOrientation4','chestOrientation1', 'chestOrientation2',
    ↳'chestOrientation3', 'chestOrientation4','ankleOrientation1',
    ↳'ankleOrientation2', 'ankleOrientation3', 'ankleOrientation4'],axis = 1)
dataCollection = dataCollection.drop(dataCollection[dataCollection.activityID
    ↳== 0].index)
dataCollection = dataCollection.apply (pd.to_numeric, errors='coerce')
dataCollection = dataCollection.dropna()

```

```

[8]: dataCollection.insert(1,'ActivityLevel',dataCollection['activityID'].
    ↳apply(map_met))

```

```

[9]: dataCollection.insert(4,'handMag',np.
    ↳sqrt((dataCollection['handMagne1']**2)+(dataCollection['handMagne2']**2)+(dataCollection['h
dataCollection.insert(5,'handAcc',np.
    ↳sqrt((dataCollection['handAcc16_1']**2)+(dataCollection['handAcc16_2']**2)+(dataCollection[
dataCollection.insert(6,'handGyro',np.
    ↳sqrt((dataCollection['handGyro1']**2)+(dataCollection['handGyro2']**2)+(dataCollection['han

```

```

dataCollection.insert(7, 'chestMag', np.
    ↳sqrt((dataCollection['chestMagne1']**2)+(dataCollection['chestMagne2']**2)+(dataCollection[
dataCollection.insert(8, 'chestAcc', np.
    ↳sqrt((dataCollection['chestAcc16_1']**2)+(dataCollection['chestAcc16_2']**2)+(dataCollection[
dataCollection.insert(9, 'chestGyro', np.
    ↳sqrt((dataCollection['chestGyro1']**2)+(dataCollection['chestGyro2']**2)+(dataCollection['c

dataCollection.insert(10, 'ankleMag', np.
    ↳sqrt((dataCollection['ankleMagne1']**2)+(dataCollection['ankleMagne2']**2)+(dataCollection[
dataCollection.insert(11, 'ankleAcc', np.
    ↳sqrt((dataCollection['ankleAcc16_1']**2)+(dataCollection['ankleAcc16_2']**2)+(dataCollection[
dataCollection.insert(12, 'ankleGyro', np.
    ↳sqrt((dataCollection['ankleGyro1']**2)+(dataCollection['ankleGyro2']**2)+(dataCollection['a

```

```

[10]: dataCollection=dataCollection.
    ↳drop(['handMagne1', 'handMagne2', 'handMagne1', 'handAcc16_1', 'handAcc16_2', 'handAcc16_3', 'hand
    ↳= 1)

```

```

[11]: dataCollection

```

```

[11]:
      timestamp ActivityLevel  activityID  heartrate  handMag  handAcc  \
2932         37.70         light          1        100.0  95.450583  9.751993
2943         37.81         light          1        100.0  95.746781  9.583806
2954         37.92         light          1        100.0  95.373006  9.639631
2965         38.03         light          1        100.0  95.360173  10.045489
2976         38.14         light          1        101.0  95.350742  9.963993
...
2871975      94.66      vigorous          24        162.0  28.812324  10.876780
2871986      94.77      vigorous          24        162.0  26.870687  10.341491
2871997      94.88      vigorous          24        162.0  26.981285  10.021582
2872007      94.98      vigorous          24        162.0  26.132942  9.932805
2872018      95.09      vigorous          24        162.0  26.239124  9.396202

      handGyro  chestMag  chestAcc  chestGyro  ankleMag  ankleAcc  \
2932    0.072467  66.439341  9.875840  0.041521  91.396062  9.956473
2943    0.435808  67.185070  9.859305  0.058315  91.797093  9.946165
2954    0.081883  66.711313  9.798303  0.072202  92.273977  9.911287
2965    0.374651  66.350301  9.929907  0.071764  92.732261  9.760667
2976    0.378423  67.355127  9.804206  0.059240  92.425175  9.888575
...
2871975  0.157022  47.894613  9.884364  0.100384  46.055915  10.063054
2871986  0.458881  47.629733  9.895226  0.162803  45.534863  9.792079
2871997  0.415535  48.081339  9.915689  0.310067  45.661446  9.809462
2872007  0.388336  47.547335  10.178812  0.322284  45.919491  9.947890
2872018  0.375698  47.415254  9.710290  0.154072  45.917820  9.967132

```

	ankleGyro	handTemperature	chestTemperature	ankleTemperature	\
2932	0.049304	30.375	32.1875	30.75	
2943	0.024230	30.375	32.1875	30.75	
2954	0.022479	30.375	32.1875	30.75	
2965	0.016725	30.375	32.1875	30.75	
2976	0.067042	30.375	32.1875	30.75	
...	
2871975	0.037724	25.125	32.3750	31.50	
2871986	0.010443	25.125	32.3750	31.50	
2871997	0.051387	25.125	32.3750	31.50	
2872007	0.042514	25.125	32.3750	31.50	
2872018	0.046552	25.125	32.3750	31.50	

	subject_id
2932	1
2943	1
2954	1
2965	1
2976	1
...	...
2871975	9
2871986	9
2871997	9
2872007	9
2872018	9

[175498 rows x 17 columns]

After the cleaning up of data we will split the dataframe into two, training set and testing set. This is mainly for the modelling which we will do soon. we call the function `train_test_split` where we will pass the dataframe, the ratio at which it need to be divided and the random state. It will return two sets which will be stored in training and testing respectively. As we can see we have a training set of 122848 samples and a testing set of 52650 samples.

```
[12]: training_data, testing_data = train_test_split(dataCollection, test_size=0.3,
    random_state=25)

print(f"No. of training examples: {training_data.shape[0]}")
print(f"No. of testing examples: {testing_data.shape[0]}")
```

No. of training examples: 122848

No. of testing examples: 52650

```
[13]: training_data
```

	timestamp	ActivityLevel	activityID	heartrate	handMag	handAcc	\
2625644	1702.08	vigorous	12	139.0	53.854060	16.163897	

1098143	224.64	light	1	73.0	19.638085	9.918705
1870506	904.86	light	17	84.0	27.798411	9.314533
1097300	216.21	light	1	72.0	19.790603	9.753240
1059712	2368.59	moderate	4	119.0	48.852229	7.863950
...
2151678	98.44	light	1	83.0	25.068682	9.776432
1524954	1196.98	light	17	88.0	4.065840	9.233299
85156	859.94	light	17	104.0	59.312579	9.262748
1504359	991.03	light	17	88.0	46.655418	13.875824
474927	990.74	light	17	88.0	51.102757	9.585142

	handGyro	chestMag	chestAcc	chestGyro	ankleMag	ankleAcc	\
2625644	0.931301	49.590104	13.070887	1.081404	43.563839	10.373201	
1098143	0.018474	49.338747	9.330957	0.058911	30.334974	9.912648	
1870506	0.463890	39.577322	9.914426	0.140350	58.405184	9.987989	
1097300	0.023966	50.097270	9.627314	0.026787	30.046044	9.877875	
1059712	1.997366	35.950529	13.262695	0.554969	44.687373	19.128132	
...	
2151678	0.032491	48.153041	9.658415	0.033238	31.660566	10.213012	
1524954	1.458080	40.232291	9.796507	0.794053	60.343038	10.061185	
85156	0.997393	57.924334	10.002432	0.676013	72.430504	9.786085	
1504359	2.601184	38.412373	9.615096	0.698069	55.510309	8.623337	
474927	1.675696	34.466511	9.707564	0.193631	55.463277	9.902596	

	ankleGyro	handTemperature	chestTemperature	ankleTemperature	\
2625644	1.537907	34.2500	38.0625	34.1875	
1098143	0.132004	32.8125	34.8125	33.5000	
1870506	0.048180	33.8125	36.1875	34.9375	
1097300	0.031313	32.8125	34.8125	33.5000	
1059712	3.563013	29.4375	36.8750	34.5000	
...	
2151678	0.046559	30.2500	32.6875	30.1250	
1524954	0.191315	35.4375	37.5625	34.5000	
85156	0.143261	33.2500	35.0000	33.7500	
1504359	0.565017	35.3125	37.3125	34.4375	
474927	0.082582	34.3125	37.6250	35.0625	

	subject_id
2625644	8
1098143	4
1870506	6
1097300	4
1059712	3
...	...
2151678	7
1524954	5
85156	1


```
1504359      5
474927       2
```

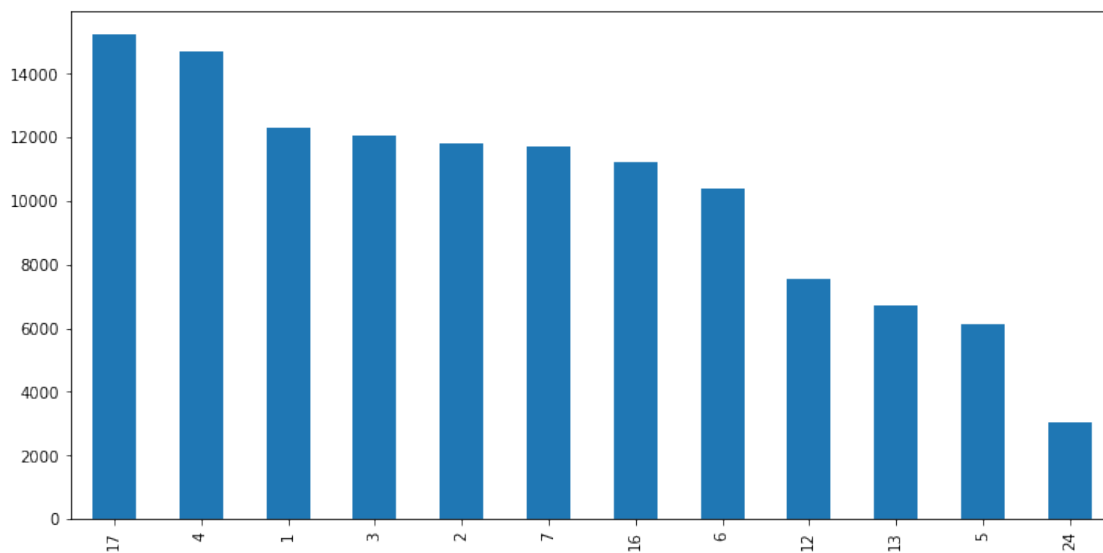
```
[122848 rows x 17 columns]
```

3 EDA

With the data partitioned into development and testing, exploratory data analysis can now be performed on our development dataset.

The first one is a bar graph where we are plotting the activity to their sample size in the dataframe. As we can see most of them are evenly distributed. Most of it are evenly balanced. only rope jumping is a bit less which is a vigorous activity.

```
[14]: training_data['activityID'].value_counts().plot(kind = "bar",figsize = (12,6))
plt.show()
```

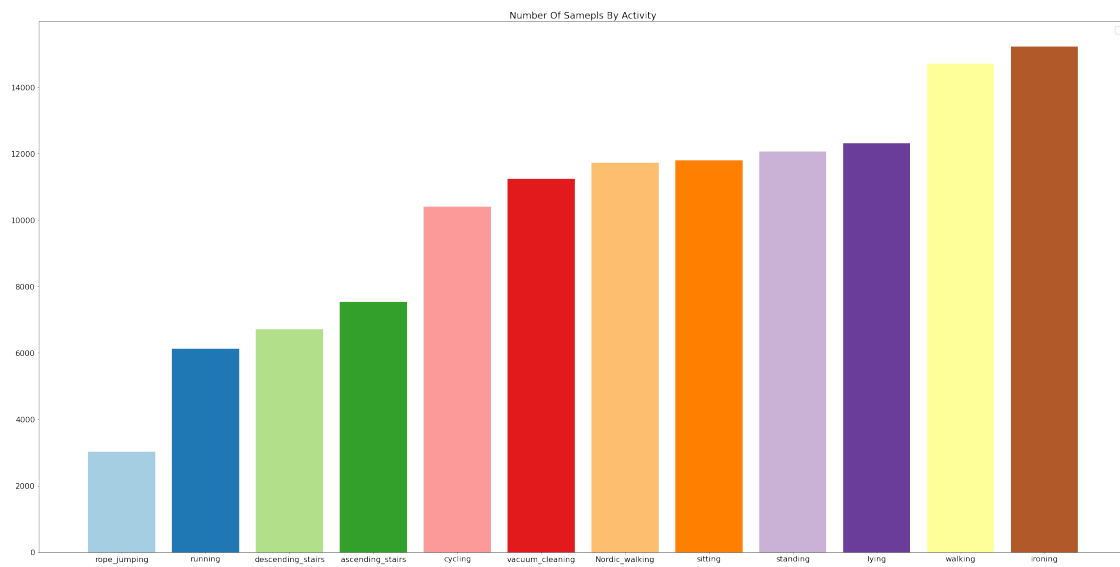


```
[15]: map_ac = load_activity_map()
sampels = training_data.groupby(['activityID']).count().reset_index()
sampels_to_subject = pd.DataFrame()
sampels_to_subject['activity'] = [map_ac[x] for x in sampels['activityID']]
sampels_to_subject['sampels'] = sampels['timestamp']
sampels_to_subject = sampels_to_subject.sort_values(by=['sampels'])
pd_fast_plot(sampels_to_subject, 'sampels', 'activity', 'Number Of Samepls By_
↳Activity',figsize=(40,20))
```

```
<ipython-input-4-fafbde2c1dde>:28: UserWarning: FixedFormatter should only be
used together with FixedLocator
```

```
a = ax.set_xticklabels(pd[column_b])
```

No handles with labels found to put in legend.



Now we are plotting the same thing but as per the activity level. As we can see below light and moderate have very large sample size compared to vigorous which is only around 11000, while the other two are greater than 50000. As defined by their MET will be looked for. The activities in the protocol can be classified as:

Light Effort (<3.0 METs): lying, sitting, standing and ironing.

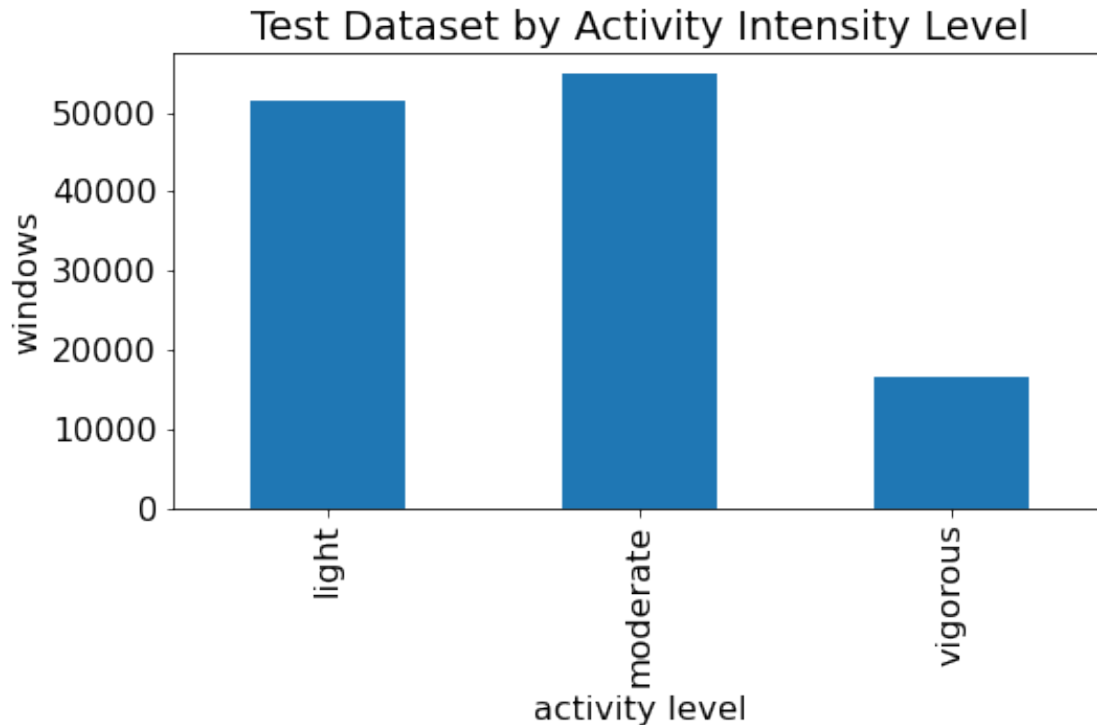
Moderate Effort (3.0 - 6.0 METs): vacuum cleaning, descending stairs, walking, Nordic walking and cycling

Vigorous Effort (> 6.0 MET): ascending stairs, running and rope jumping

```
[16]: s = training_data.groupby('ActivityLevel').count()['activityID']
s = s.rename("Activity Level Counts")
print(('Dev Dataset by Activity Intensity Level'))
display(s)
ax = s.plot(kind='bar', figsize=(8,4))
_ = ax.set_ylabel('windows')
_ = ax.set_xlabel('activity level')
_ = ax.set_title('Test Dataset by Activity Intensity Level')
```

Dev Dataset by Activity Intensity Level

```
ActivityLevel
light      51390
moderate   54787
vigorous   16671
Name: Activity Level Counts, dtype: int64
```



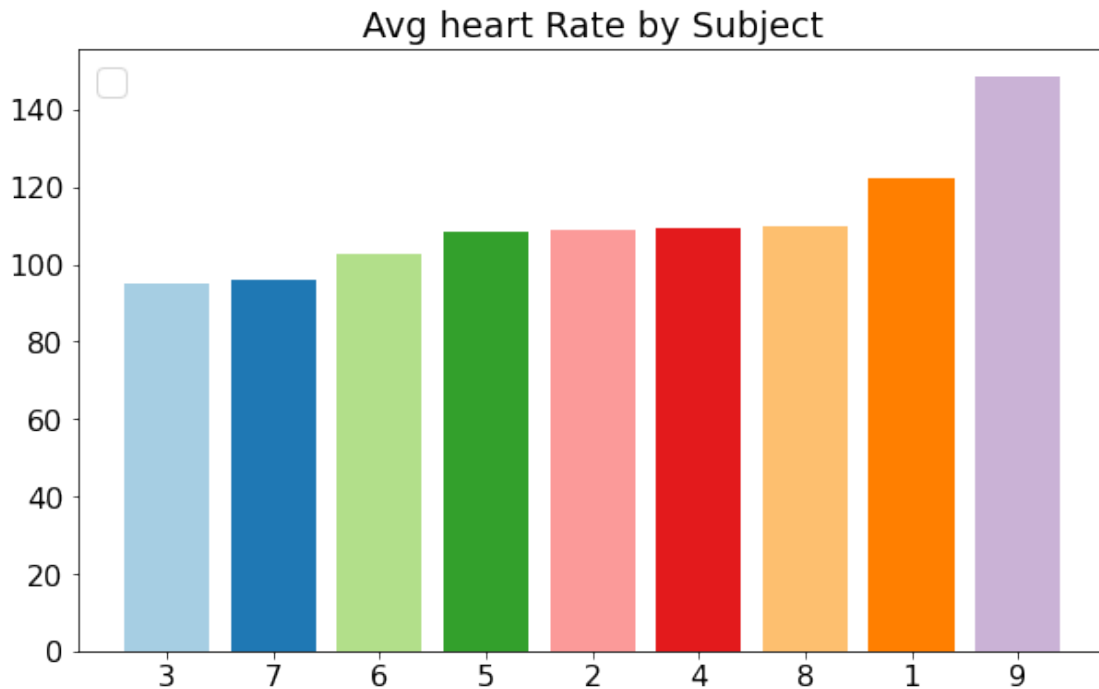
For the next one we will plot the subject IDs with their average heart rate. For the same we will be using a bar graph. A data frame result id is created where we take the data from training_ data and it is grouped by subject id. From the graph we can see that the subject 9 has mch higher hear rate than any other subject. Its above 140 while the rest of them are between 90 and 120. We are going to focus on heart rate as it is our most precice meter of check for tracking subjects during activities as implied by the various indications on the readme file of the dataset.

```
[17]: result_id = training_data.groupby(['subject_id']).mean().reset_index()

sampels_heart_rate = pd.DataFrame()
sampels_heart_rate['id'] = result_id['subject_id']
sampels_heart_rate['heartrate'] = result_id['heartrate']
sampels_heart_rate = sampels_heart_rate.sort_values(by=['heartrate'])
pd_fast_plot(sampels_heart_rate, 'heartrate', 'id', 'Avg heart Rate by Subject')
```

<ipython-input-4-fafbde2c1dde>:28: UserWarning: FixedFormatter should only be used together with FixedLocator

```
a = ax.set_xticklabels(pd[column_b])
No handles with labels found to put in legend.
```



The next graph is also a bar graph with the activity levels against the average heart rates. For this we create a new data frame first, `ha` in which we are grouping by activity level and taking the mean of heart rate. Then we are using this `ha` data frame to plot the graph. From the graph we can infer that the heart rate of light activities is less compared to moderate and the heart rate of moderate activities is less compared to vigorous.

```
[18]: ha = training_data.groupby('ActivityLevel').mean()['heartrate']

print(('Activity heart rate'))
display(ha)
ax = ha.plot(kind='bar', figsize=(8,4))
_ = ax.set_ylabel('Heart Rate')
_ = ax.set_xlabel('activity level')
_ = ax.set_title('Heartrate of Activity Levels')
```

Activity heart rate

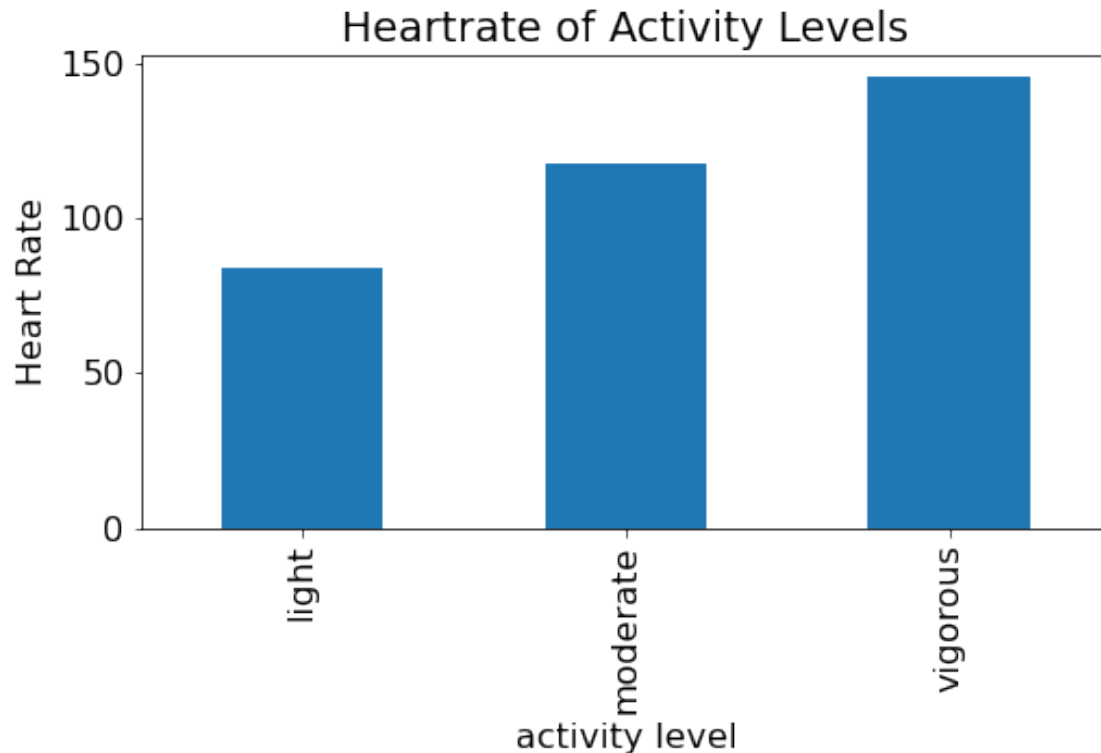
ActivityLevel

light 83.888266

moderate 117.606586

vigorous 145.141623

Name: heartrate, dtype: float64



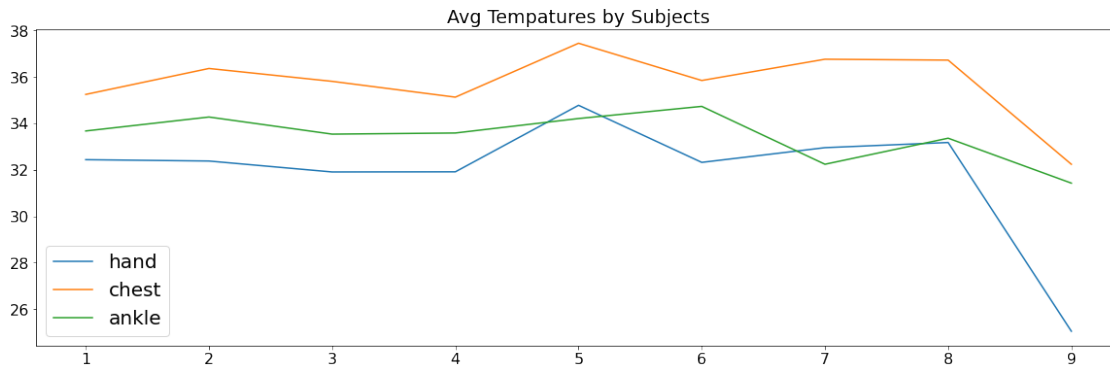
For the next graph we are plotting the average temperature by subject. For this we are taking the temperature of hand, chest and ankle. First we create a dataframe `samepls_tempreture`. Then we put the values of hand temperature, chest temperature and ankle temperature into it from the data frame `result_id`. Once we do that we will pass that to the plot function and get the graph. From the graph we can see that the chest temperature is high for all the subjects followed by ankle and hand. But when it comes to 9 everything falls so low even though the average heartbeat was higher or 9. This must be an issue in the data.

```
[19]: samepls_tempreture = pd.DataFrame()
samepls_tempreture['hand'] = result_id['handTemperature']
samepls_tempreture['chest'] = result_id['chestTemperature']
samepls_tempreture['ankle'] = result_id['ankleTemperature']

ax = samepls_tempreture.plot(kind='line', figsize=(20,6), title='Avg Temperatures_
↳by Subjects')
a = ax.set_xticklabels(result_id['subject_id'])
b = ax.legend(fontsize = 20)
c = ax.set_xticks(np.arange(len(samepls_tempreture)))
```

<ipython-input-19-e946056f7bd5>:7: UserWarning: FixedFormatter should only be used together with FixedLocator

```
a = ax.set_xticklabels(result_id['subject_id'])
```

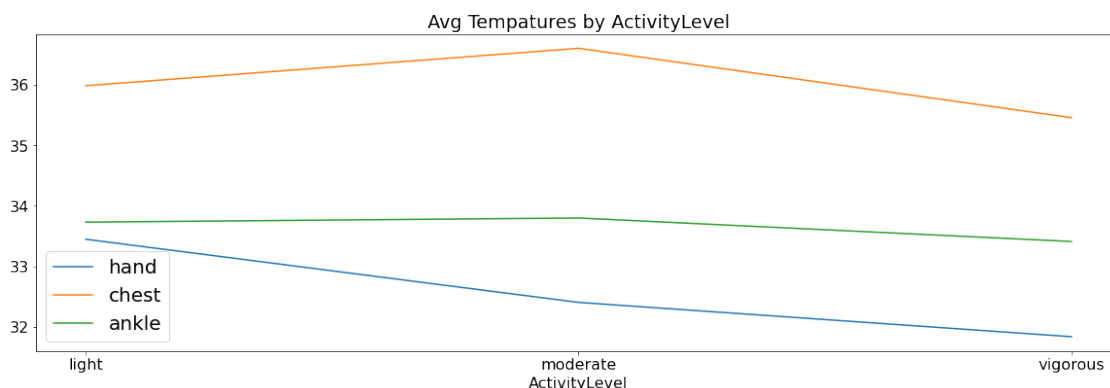


A new data frame eta is created where we are grouping it by activity level. Then we are creating dataframe sample temperature like we did above to take the temperature for hand chest and ankle. Once thats done we will use it to plot the activity level against the temperature. And from the graph we can see that the chest temperature is still the highest followed by ankle and hand, but then again the temperature for vigorous activity level is still the loest which shouldnt be the case.

```
[20]: ta = training_data.groupby('ActivityLevel').mean()
```

```
[21]: samepls_tempreture = pd.DataFrame()
samepls_tempreture['hand'] = ta['handTemperature']
samepls_tempreture['chest'] = ta['chestTemperature']
samepls_tempreture['ankle'] = ta['ankleTemperature']

ax = samepls_tempreture.plot(kind='line', figsize=(20,6), title='Avg Temperatures_
↳by ActivityLevel')
#a = ax.set_xticklabels(ta['ActivityLevel'])
b = ax.legend(fontsize = 20)
c = ax.set_xticks(np.arange(len(samepls_tempreture)))
```

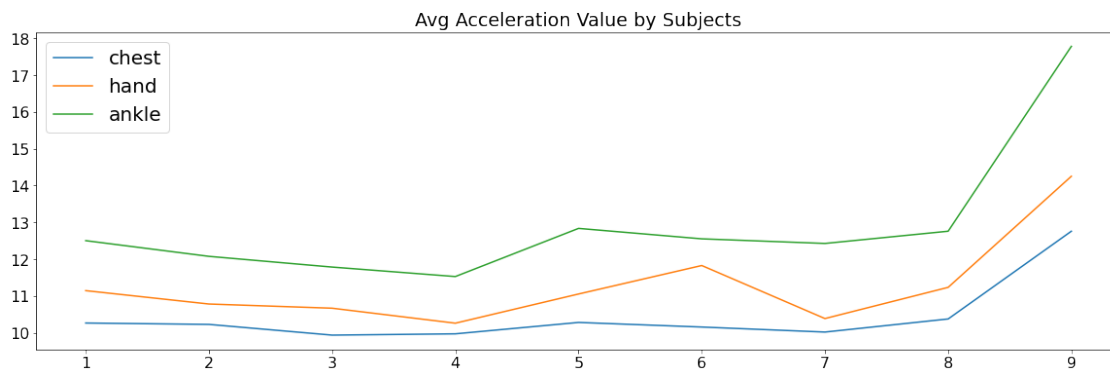


In the next graph we are plotting the acceleration of chest, hand and ankle of each subjects. For this we will use the chestacc , handacc and ankleacc calculated in the beginning using the three points in the 3d space. while creating the dataframe result_id we have taken the mean of those and have grouped by subject ids, we use that for this. Like above we create a new data frame samepls where we put chestacc hand acc and ankle acc. Then we pass it for plotting. From the linegraph plotted we can see that the ankle has the most acceleration than any other body part, which is then followed by the hand and then the chest. The acceleration is almost the same for all except 9 where it goes really high.

```
[22]: samepls = pd.DataFrame()
samepls['chest'] = result_id['chestAcc']
samepls['hand'] = result_id['handAcc']
samepls['ankle'] = result_id['ankleAcc']
ax = samepls.plot(kind='line', figsize=(20,6), title='Avg Acceleration Value by_
↳Subjects')
a = ax.set_xticklabels(result_id['subject_id'])
b = ax.legend(fontsize = 20)
c = ax.set_xticks(np.arange(len(samepls)))
```

<ipython-input-22-fc9e94688e69>:6: UserWarning: FixedFormatter should only be used together with FixedLocator

```
a = ax.set_xticklabels(result_id['subject_id'])
```



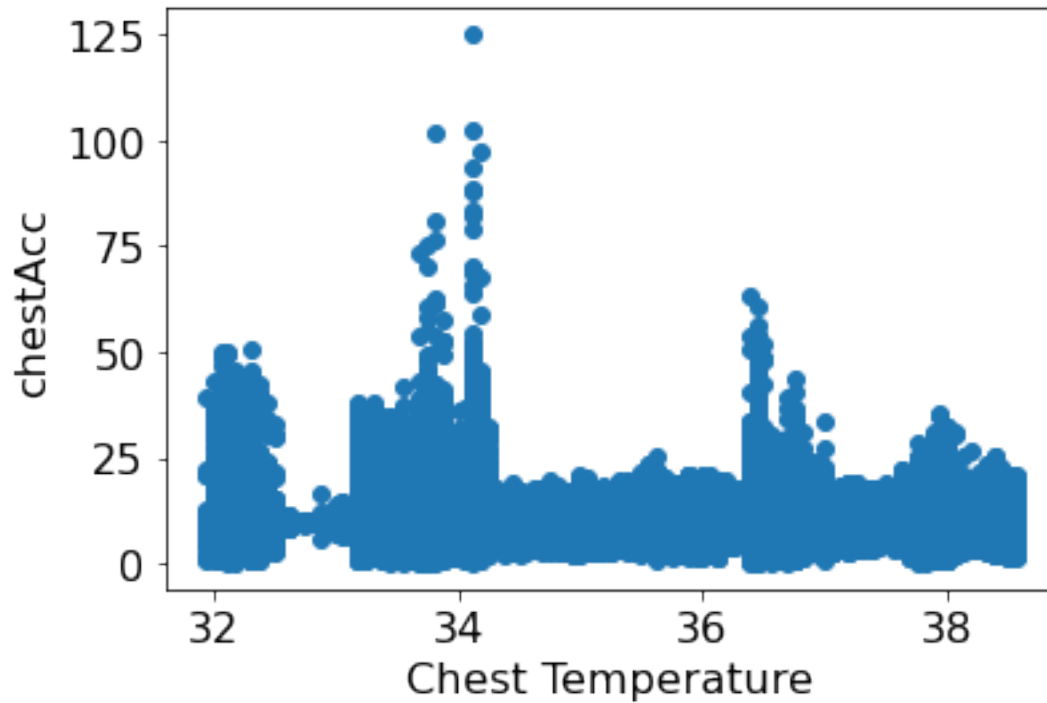
Now we are plotting the Hand Temperature against Hand Acceleration, chest temperature against chest acceleration and ankle temperature against ankle acceleration. For this we have used scatter plot and we can see that its mostly distributed. the chest acceleration is almost the same for all temperature. Likewise for the rest too.

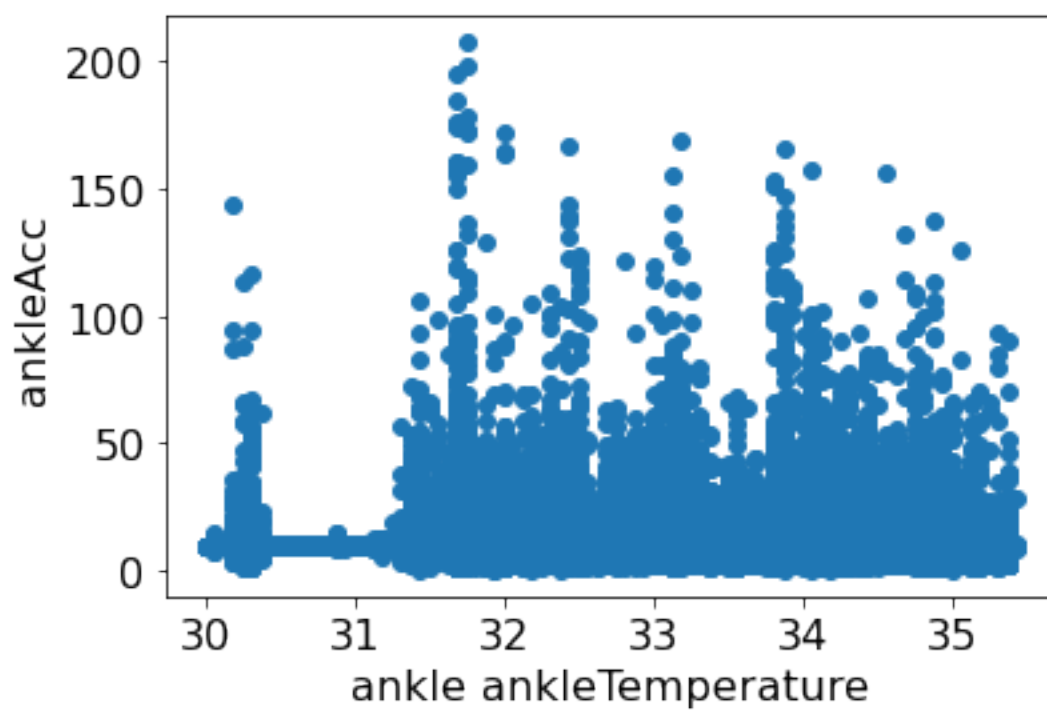
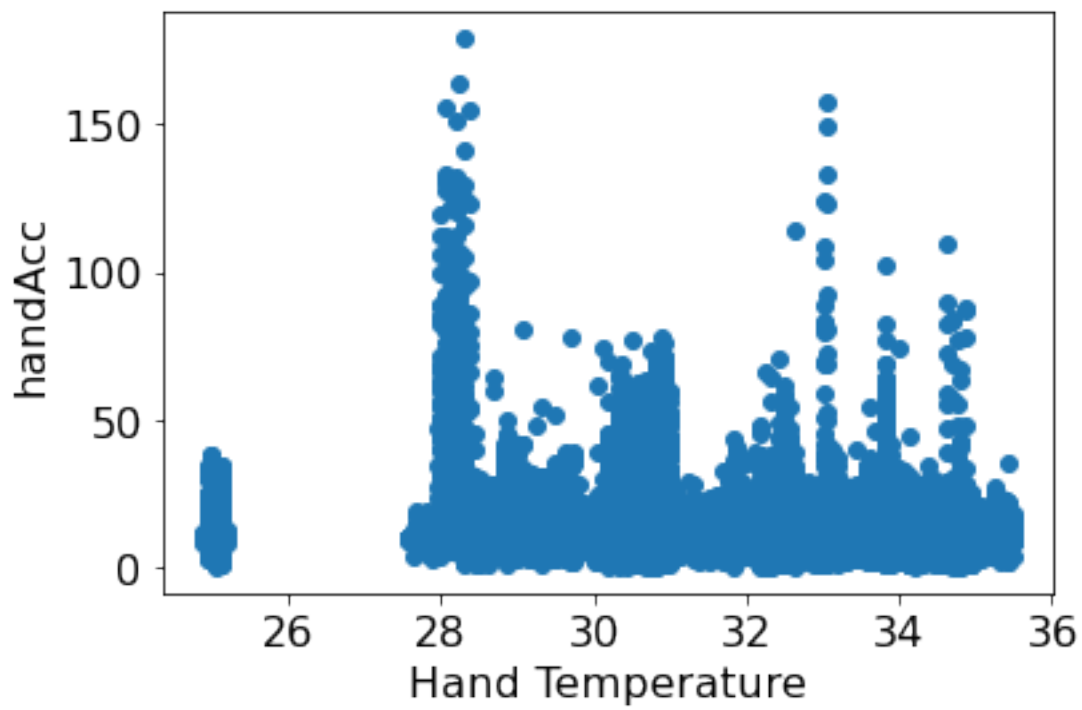
```
[390]: plt.scatter(training_data['chestTemperature'], training_data['chestAcc'])
plt.xlabel('Chest Temperature')
plt.ylabel('chestAcc')
plt.show()

plt.scatter(training_data['handTemperature'], training_data['handAcc'])
plt.xlabel('Hand Temperature')
```

```
plt.ylabel('handAcc')  
plt.show()
```

```
plt.scatter(training_data['ankleTemperature'], training_data['ankleAcc'] )  
plt.xlabel('ankle ankleTemperature')  
plt.ylabel('ankleAcc')  
plt.show()
```





4 Hypothesis

During EDA, the Heart Rate was identified as having the potential to allow classification of activities according to their MET equivalent. To check this a set two two hypothesis will be created which will be using the mean of heart rate for each of the MET levels. The mean of the heart rate according to the graphs we got above should be like Light is less than moderate which is less than vigorous.

The First hypothesis is with the light activities and the moderate activities.

4.0.1 1: If Heartrate is related with Activity level, then Heart rate of moderate activity level is greater than Heartrate of light activity level.

H0: The mean heartrate of moderate activities have no difference with light activities

H1: The mean heartrate of moderate activities is higher than that of light activities

For the hypothesis we are creating three new data frames, one for each activity level. Then in those data frames we are only putting data corresponding to that activity level

```
[35]: lightdf=testing_data[testing_data['ActivityLevel']=='light']
      moddf=testing_data[testing_data['ActivityLevel']=='moderate']
      vigdf=testing_data[testing_data['ActivityLevel']=='vigorous']
```

After that we calculate the value of p value of difference. for that we have to subtract the mean heartrate of moderate and light and then divide it by the squareroot of (moderate standard deviation/count)² -(light standard deviation/count)² . Then we will get the a value which will be stored in pv. Then the variable pv will be used to get the p value, where it will be passed to the scipy function norm.cdf which will return a value, it will be stored in Pvalue.

```
[36]: pv=((moddf['heartrate'].mean())-(lightdf['heartrate'].mean()))/
      ↪((moddf['heartrate'].std()**2)/(moddf['heartrate'].
      ↪count()**2))-((lightdf['heartrate'].std()**2)/(lightdf['heartrate'].
      ↪count()**2))
```

```
[37]: import scipy.stats
      pValue = 1 - scipy.stats.norm.cdf(pv)
      pValue
```

```
[37]: 0.0
```

With a P value of 0, the null hypothesis can be rejected. As such it can be concluded that Moderate effort activity levels will have a higher mean HR than light effort activity levels.

The Second hypothesis is with the moderate activities and the vigorous activities. ### 2: If Heartrate is related with Activity level, then Heart rate of vigorous activity level is greater than Heartrate of moderate activity level.

H0: The mean heartrate of vigorous activities have no difference with moderate activities

H1: The mean heartrate of vigorous activities is higher than that of moderate activities

We are calculating the P value the same way we did above. We have to subtract the mean heartrate of vigorous and moderate and then divide it by the squareroot of $(\text{vigorous standard deviation}/\text{count})^2 - (\text{moderate standard deviation}/\text{count})^2$. Then we will get the a value which will be stored in pv. Then the variable pv will be used to get the p value, where it will be passed to the scipy function norm.cdf which will return a value, it will be stored in Pvalue.

```
[38]: pv=((vigdf['heartrate'].mean())-(moddf['heartrate'].mean()))/
      ↪(((vigdf['heartrate'].std()**2)/(vigdf['heartrate'].
      ↪count()**2))-((moddf['heartrate'].std()**2)/(moddf['heartrate'].count()**2)))
```

```
[39]: import scipy.stats
      pValue = 1 - scipy.stats.norm.cdf(pv)
      pValue
```

```
[39]: 0.0
```

With the p value coming as 0.0, we can say that the null hypothesis can be rejected. Hence it can be concluded that the vigorous activity level will have higher mean HR than moderate activity level.

From the above two hypothesis tests, we can conclude that heartrate and MET value will be enough to identify the activities in the dataset.

4.0.2 3: We want to see if there's any relation between heartrate and temperature.

H0: Chest Temperature is not dependent on Heartrate

H1: Chest Temperature is dependent on Heartrate

For the same we are using the pearsons correlation method. And we calculate the regression r value and p value to make the decision.

```
[26]: from scipy.stats import pearsonr

      d1=testing_data['heartrate']
      d2=testing_data['chestTemperature']
      stat, p = pearsonr(d1, d2)
      print('stat= %.3f, p= %.3f' % (stat, p))
```

```
stat=-0.120, p=0.000
```

Since the p-value is less than 0.05, the null hypothesis can be rejected. Hence from H1, we can conclude that the Chest temperature is dependent on heart rate.

5 Modelling

For modelling the first thing is to create a training set without any labels like activityID, timestamp, subjectID etc. After that we are creating another dataframe train1 which has activityID and subject ID, this will be used afterwards for creating a dataframe.

```
[330]: training_data_temp=training_data.
        ↪drop(['activityID','timestamp','subject_id','handGyro','chestGyro','ankleGyro','chestMag',''])
        train1 = training_data[['activityID','subject_id']]
```

```
[331]: training_data_temp
```

```
[331]:
```

	heartrate	handAcc	chestAcc	ankleAcc	handTemperature \
2625644	139.0	16.163897	13.070887	10.373201	34.2500
1098143	73.0	9.918705	9.330957	9.912648	32.8125
1870506	84.0	9.314533	9.914426	9.987989	33.8125
1097300	72.0	9.753240	9.627314	9.877875	32.8125
1059712	119.0	7.863950	13.262695	19.128132	29.4375
...
2151678	83.0	9.776432	9.658415	10.213012	30.2500
1524954	88.0	9.233299	9.796507	10.061185	35.4375
85156	104.0	9.262748	10.002432	9.786085	33.2500
1504359	88.0	13.875824	9.615096	8.623337	35.3125
474927	88.0	9.585142	9.707564	9.902596	34.3125

	chestTemperature	ankleTemperature
2625644	38.0625	34.1875
1098143	34.8125	33.5000
1870506	36.1875	34.9375
1097300	34.8125	33.5000
1059712	36.8750	34.5000
...
2151678	32.6875	30.1250
1524954	37.5625	34.5000
85156	35.0000	33.7500
1504359	37.3125	34.4375
474927	37.6250	35.0625

[122848 rows x 7 columns]

train_data1 is created where training_data is converted to array. Then we do the clustering which is the process of grouping data in such a way that objects in the same group or cluster are more similar to each other than to those in other clusters. We use the K-means clustering in which the similarity between cluster points is derived by the closeness of a point to the centroid of the cluster.

```
[332]: train_data1=training_data_temp.to_numpy()

        K=3 # specify number of clusters
        #Clustering
        cl_K3=cluster.KMeans(init='random',n_clusters=K)
        cl_K3.fit(train_data1)
        cl_K3.predict(train_data1)
```

```
[332]: array([0, 1, 1, ..., 0, 1, 1])
```

Then we use the predict function to predict the label for all the data and it will be stored to ax.

```
[345]: ax = cl_K3.predict(training_data_temp)
       print(ax)
```

```
[0 1 1 ... 0 1 1]
```

A data frame train_Data1 is created, where we put a copy of training_Data_temp and then add two more columns, one for activity and the other for predicted label which we created above.

```
[346]: train_data1=training_data_temp.copy(deep=True)
       train_data1['activity']=train1['activityID']
       train_data1['predicted_label']=ax
       train_data1.head(10)
```

```
#map_ac[x] for x in samples['activityID']
```

```
[346]:
```

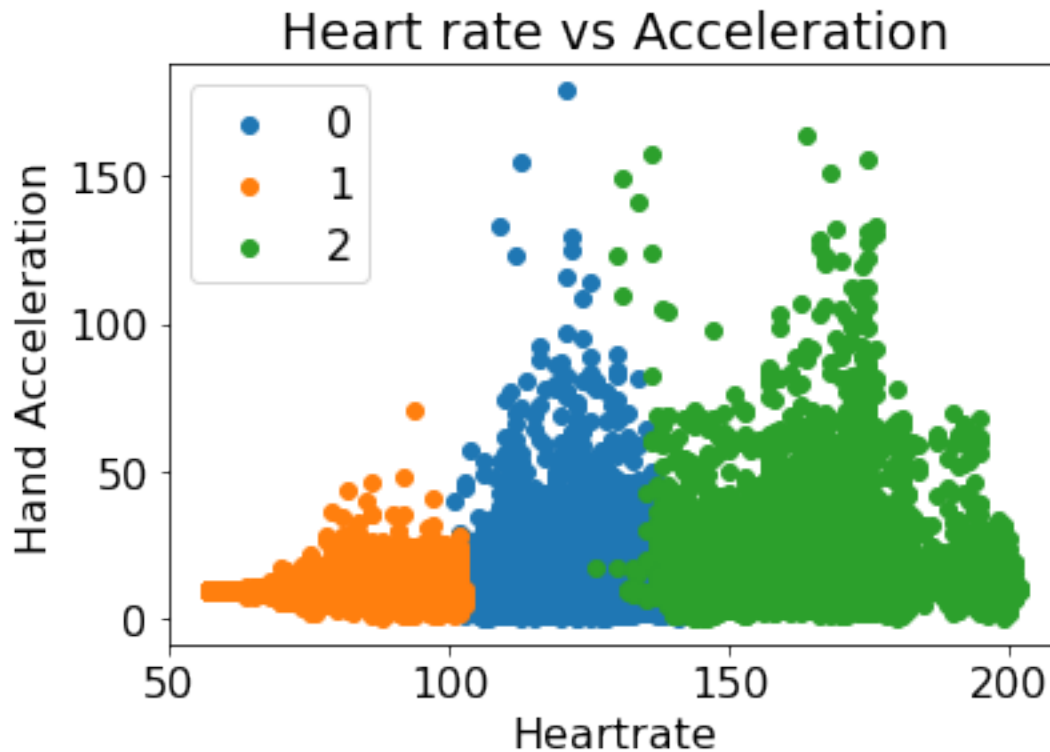
	heartrate	handAcc	chestAcc	ankleAcc	handTemperature	\
2625644	139.0	16.163897	13.070887	10.373201	34.2500	
1098143	73.0	9.918705	9.330957	9.912648	32.8125	
1870506	84.0	9.314533	9.914426	9.987989	33.8125	
1097300	72.0	9.753240	9.627314	9.877875	32.8125	
1059712	119.0	7.863950	13.262695	19.128132	29.4375	
2229737	84.0	9.289872	10.252639	9.862815	33.5625	
2030534	94.0	6.605565	6.516699	10.705343	31.7500	
1723120	133.0	28.771755	12.435896	15.497875	34.1250	
2524767	80.0	9.729208	9.716566	9.935296	34.4375	
1201439	107.0	8.067283	8.434496	10.075797	33.6875	

	chestTemperature	ankleTemperature	activity	predicted_label
2625644	38.0625	34.1875	12	0
1098143	34.8125	33.5000	1	1
1870506	36.1875	34.9375	17	1
1097300	34.8125	33.5000	1	1
1059712	36.8750	34.5000	4	0
2229737	36.3750	32.6875	17	1
2030534	36.5000	34.8750	4	1
1723120	37.6250	34.1875	6	0
2524767	37.5625	34.1875	3	1
1201439	36.0000	34.4375	16	0

We plot the Heartrate against hand acceleration to see how the clusters are created, and as we can see from the graph below, its divided into three clusters, the orange one is the first one, the blue the second and green the third.

```
[356]: plt.scatter(train_data1[train_data1['predicted_label']==0] .
               ↳heartrate,train_data1[train_data1['predicted_label']==0] .handAcc)
```

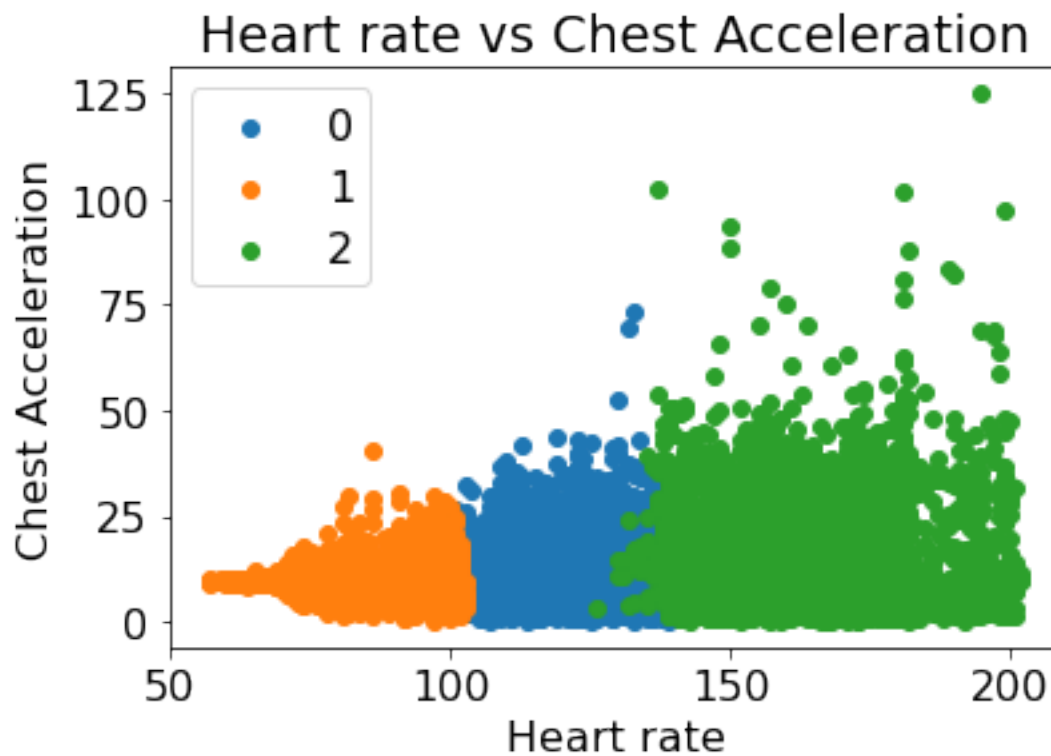
```
plt.scatter(train_data1[train_data1['predicted_label']==1].
    ↳heartrate,train_data1[train_data1['predicted_label']==1].handAcc)
plt.scatter(train_data1[train_data1['predicted_label']==2].
    ↳heartrate,train_data1[train_data1['predicted_label']==2].handAcc)
plt.xlabel('Heartrate')
plt.ylabel('Hand Acceleration')
plt.legend([0,1,2])
plt.title('Heart rate vs Acceleration')
plt.show()
```



The same way we plot graphs for Heart Rate vs Chest Acceleration, from 50 to 100 hear rate it is one clcluster, then from 100 to 150 another one and 150 to 200 third one.

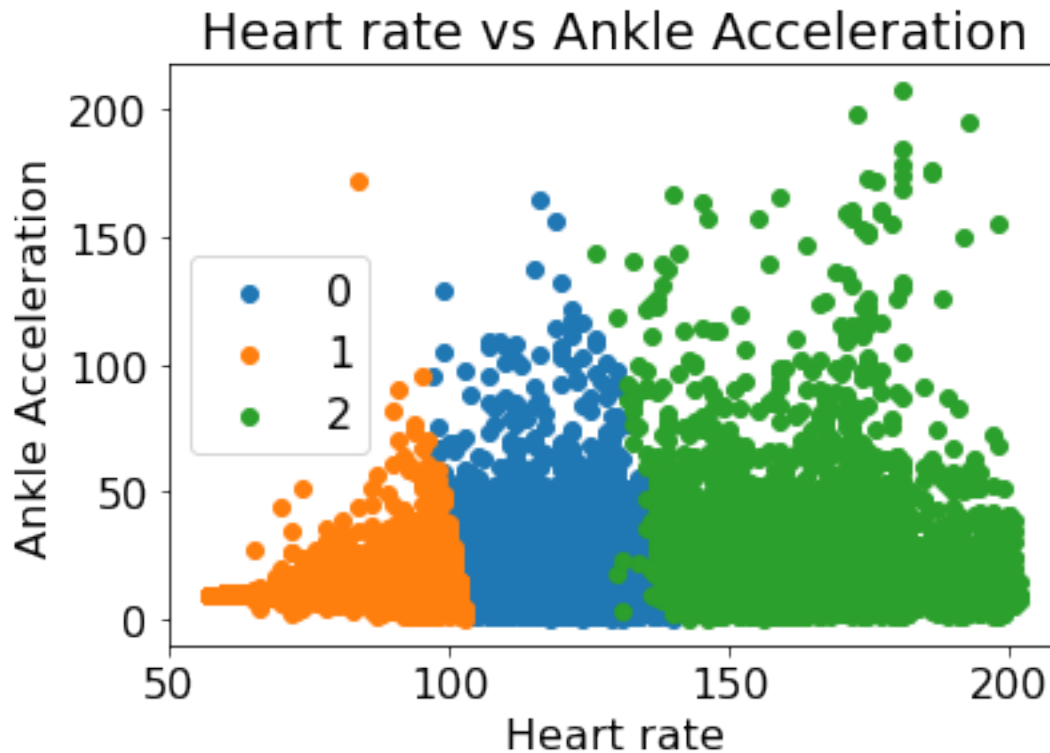
```
[358]: plt.scatter(train_data1[train_data1['predicted_label']==0].
    ↳heartrate,train_data1[train_data1['predicted_label']==0].chestAcc)
plt.scatter(train_data1[train_data1['predicted_label']==1].
    ↳heartrate,train_data1[train_data1['predicted_label']==1].chestAcc)
plt.scatter(train_data1[train_data1['predicted_label']==2].
    ↳heartrate,train_data1[train_data1['predicted_label']==2].chestAcc)
plt.xlabel('Heart rate')
plt.ylabel('Chest Acceleration')
plt.legend([0,1,2])
```

```
plt.title('Heart rate vs Chest Acceleration')
plt.show()
```



The same way we plot graphs for Heart Rate vs ankle Acceleration, from 50 to 100 hear rate it is one clcluster, then from 100 to 150 another one and 150 to 200 third one.

```
[359]: plt.scatter(train_data1[train_data1['predicted_label']==0] .
    ↳heartrate,train_data1[train_data1['predicted_label']==0].ankleAcc)
plt.scatter(train_data1[train_data1['predicted_label']==1] .
    ↳heartrate,train_data1[train_data1['predicted_label']==1].ankleAcc)
plt.scatter(train_data1[train_data1['predicted_label']==2] .
    ↳heartrate,train_data1[train_data1['predicted_label']==2].ankleAcc)
plt.xlabel('Heart rate')
plt.ylabel('Ankle Acceleration')
plt.legend([0,1,2])
plt.title('Heart rate vs Ankle Acceleration')
plt.show()
```



We create a dataframe `df_label` to see what has been predicted. For that we are taking the activity and the predicted labels and grouping by it. Each label represents the level of activities. The cluster 0 consists of Moderate activities, cluster 1 consists of Light activities and 2 consists of vigorous activities. And as we can see the activities are divided into the right labels. Activity 1,2,3 are light activities which are in label 1. Then the activity 4, which is walking is a moderate activity which is in label 2, then activity 5, Running is a vigorous activity is in label 2 and so on.

```
[338]: df_label = train_data1.groupby(['activity', 'predicted_label']).predicted_label.  
        ↪count().unstack()  
df_label
```

```
[338]: predicted_label      0      1      2  
activity  
1          335.0  11981.0    NaN  
2          207.0  11589.0    NaN  
3         1331.0  10726.0    NaN  
4        11829.0   2886.0    NaN  
5          1263.0    89.0  4772.0  
6        10199.0   135.0   66.0  
7        10456.0   326.0  940.0  
12         4223.0   763.0 2542.0  
13         3562.0   797.0 2346.0  
16         5051.0  6194.0    NaN
```


17	1758.0	13463.0	NaN
24	627.0	NaN	2392.0

Now we have to test it with the testing data, and for that we are preparing the testing set. For that we do the same thing we did for the training data. Dropping all the unnecessary columns like activityID, timestamp etc. Then we create test1 where we have two columns activityID and subjectID.

```
[362]: test_data_temp=testing_data.
      ↳drop(['activityID','timestamp','subject_id','handGyro','chestGyro','ankleGyro','chestMag','...'])
      test1 = testing_data[['activityID','subject_id']]
```

created ax test for all the predicted labels. we are passing the test_data_temp to predict the labels.

```
[341]: ax_test =cl_K3.predict(test_data_temp)
      print(ax_test)
```

```
[1 0 1 ... 1 0 1]
```

We then add two columns predicted_label and activity to the test_Data1 from the test1 data frame

```
[342]: test_data_temp['predicted_label']=ax_test
      test_data1=test_data_temp.copy(deep=True)
      test_data1['activity']=test1['activityID']
      test_data1.head(10)
```

```
[342]:      heartrate    handAcc    chestAcc    ankleAcc    handTemperature \
963970      94.0    11.408266    9.751466    10.018422      33.1875
1054560    119.0    15.373321    8.965425    10.046451      29.8750
1832564     72.0     9.789977    9.935780    10.003130      33.5000
1720189    124.0    13.897841    9.355981    16.588538      34.2500
2166356     61.0     9.693993    9.858129     9.940656      31.0000
1480592     87.0     9.831521    9.895031    10.318878      35.1250
711694     131.0     9.945269    4.309880     9.365050      30.4375
2287990     72.0    10.057362    9.727764     9.911510      34.1875
555898     144.0    13.818170    14.614707    30.361028      34.1250
1831306     72.0     9.679798    9.781048    10.180866      33.5000
```

	chestTemperature	ankleTemperature	predicted_label	activity
963970	36.6875	34.0625	1	16
1054560	36.9375	34.5625	0	4
1832564	35.8750	34.7500	1	2
1720189	37.8125	34.2500	0	6
2166356	33.8125	30.8125	1	1
1480592	37.0625	34.3125	1	3
711694	35.4375	34.1250	0	7
2287990	37.6250	33.0625	1	12
555898	38.0625	35.3125	2	12

1831306	35.8750	34.7500	1	2
---------	---------	---------	---	---

we then create a dataframe `d_test_label` which has activity and predicted labels. And we can see the activities are divided into clusters the same way it was did for the training set. Activities 1,2 and 3 are in label q which is light. Activity 4 is in label 0 which is moderate. Activity 5 is in 2 which is vigorous. Hence we can say it is working as expected

```
[343]: df_label_test = test_data1.groupby(['activity','predicted_label']).
        ↪predicted_label.count().unstack()
df_label_test
```

```
[343]: predicted_label      0      1      2
activity
1          162.0  5097.0    NaN
2           89.0  4996.0    NaN
3          612.0  4587.0    NaN
4         5060.0  1223.0    NaN
5          520.0   20.0  2001.0
6         4445.0   54.0   34.0
7         4601.0  127.0  407.0
12         1804.0  301.0 1066.0
13         1514.0  334.0 1032.0
16         2159.0 2593.0    NaN
17          733.0 5742.0    NaN
24          262.0    NaN 1075.0
```

Now we need to find the Silhouette Coefficient or silhouette score which is a metric used to calculate the goodness of a clustering technique. Its value ranges from -1 to 1.

For that we import the `silhouette_score` function from `sklearn.metrics.cluster`. Then we pass the `training_data_temp` to that function. And we we can see the score came out as 0.5205 which means the clusters are apart from each other and distinguished well.

```
[368]: from sklearn.metrics.cluster import silhouette_score
SC_3=silhouette_score(training_data,c1_K3.labels_)
print('Silhouette Score = ', SC_3)
```

```
Silhouette Score = 0.2702782813219407
```

6 Conclusion

For this report we had three objectives. First one was Cleaning up the data and exploratory data analysis. There we had removed all the transient activities and removed all the unnecessary columns as part of cleanup, then for EDA we mainly used heartrate for all the plottings and as we could see vigorous activities has more heartrate. But then we came to know that subje 9 has some issues with the attributes in the data.

Then for the second objective, we did three hypothesis. The first two was regrading the hearrate and the activity level, where we came to the conclusion that hearrate of different levels goes in the

order vigorous > moderate > light. For the third hypothesis we checked if temperature is depended on hearrate came to the conclusion that it is.

Then for the third objective modelling, we created clusters and then used the training data to see if the predictions was right. Afterwards we used the testing data set which also yielded the same results as the training set. Hence the modelling was successful.