

```
mirror_mod = modifier_ob.  
set mirror object to mirror.  
mirror_mod.mirror_object  
operation == "MIRROR_X":  
mirror_mod.use_x = True  
mirror_mod.use_y = False  
mirror_mod.use_z = False  
operation == "MIRROR_Y":  
mirror_mod.use_x = False  
mirror_mod.use_y = True  
mirror_mod.use_z = False  
operation == "MIRROR_Z":  
mirror_mod.use_x = False  
mirror_mod.use_y = False  
mirror_mod.use_z = True  
  
selection at the end -add  
mirror_ob.select= 1  
modifier_ob.select=1  
context.scene.objects.active  
("Selected" + str(modifier_ob.  
mirror_ob.select = 0  
= bpy.context.selected_object  
data.objects[one.name].select  
  
print("please select exactly  
  
-- OPERATOR CLASSES ----  
  
types.Operator):  
X mirror to the selected  
object.mirror_mirror_x"  
mirror X"  
  
context):  
context.active_object is not
```

Python Project

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1) Data-Visualization

- First of all, here are the librairies that we used to study this dataset.
- We renamed the columns of the dataset so it will be easier to manipulate them.



```
import seaborn as sns # used for plot interactive graph.  
import numpy as np # linear algebra  
import pandas as pd # data manipulation  
import matplotlib.pyplot as plt # this is used for plot and graph  
import seaborn as sns # used for plot interactive graph.
```

[199]

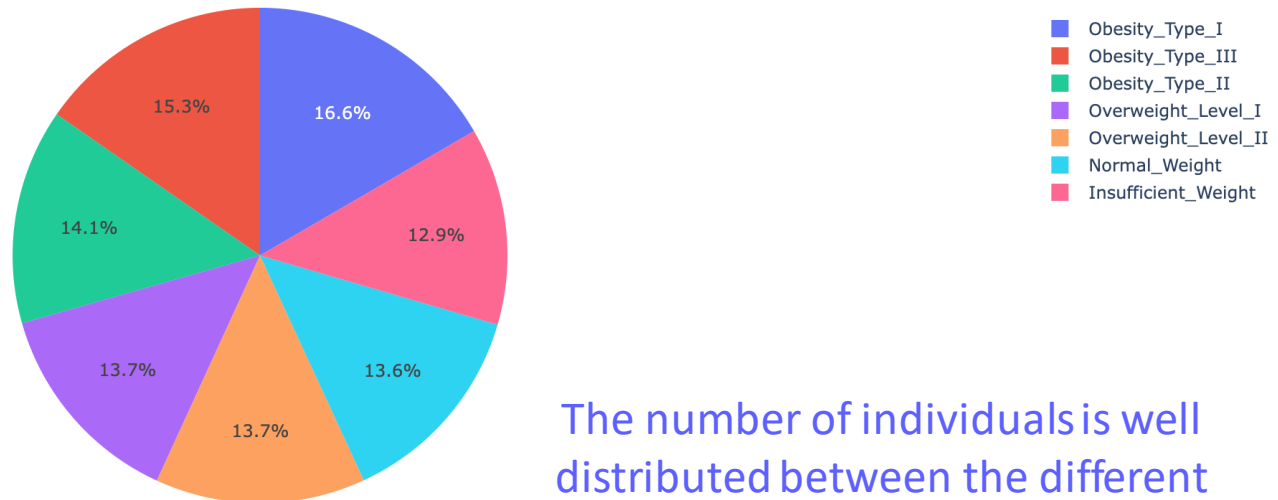
```
df=pd.read_csv("ObesityDataSet_raw_and_data_sinthetic.csv")  
df.columns=["Gender", "Age", "Height", "Weight", "Family_history",  
            "High_caloric_food", "Vegetables", "number_meals", "Snacking",  
            "Smoke", "Daily_water", "Count_calories", "Physical_activity",  
            "Hours_screen", "Alcool", "Traveling_way", "Shape"]
```

First visualization of the dataset

Our dataset is an estimation of obesity levels based on different parameters.

```
[105] fig = px.pie(df, names='Shape', title="Nombre d'individus par catégorie de poids")  
fig.show()
```

Nombre d'individus par catégorie de poids



The number of individuals is well distributed between the different morphologies

Here we will try to visualize the BMI aspect on a graph with weight in function of height.
This has nothing to see with the data from dataset, but it will help for our analysis

```
[229] def pointsimc(x, imc): #Fonction used to have the coordinates of the IMC limits
      y=imc*x*x
      return (x,y)
```



```
f, ax = plt.subplots(figsize=(8, 6))
```

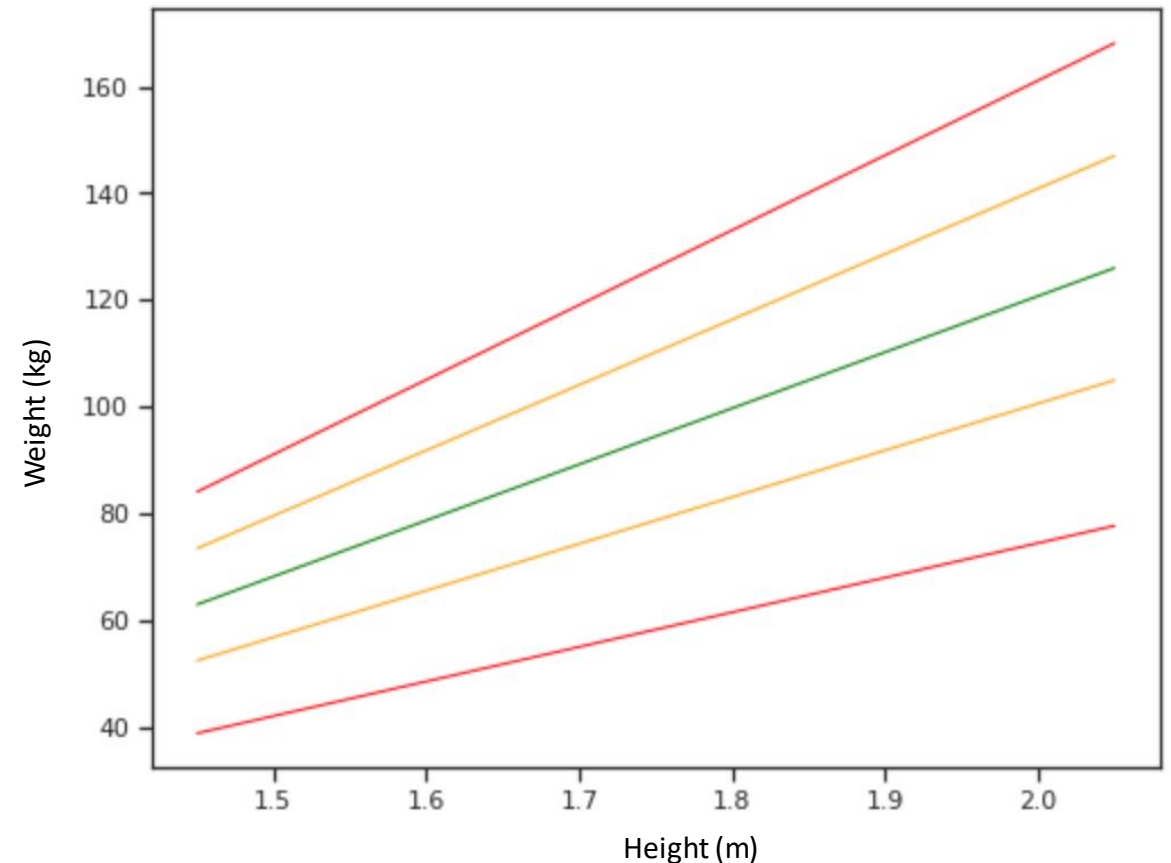
```
xunderweight=[1.45,2.05] #Min and Max heights in the dataset
yunderweight=[38.89,77.746] #calculated coordinates for each cathegory of BMI
figure1= plt.plot(xunderweight, yunderweight, c="red", linewidth =1)
```

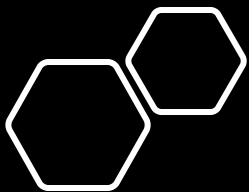
```
xnormal=[1.45,2.05] #Min and Max heights in the dataset
ynormal=[52.5,105] #calculated coordinates for each cathegory of BMI
figure1= plt.plot(xnormal, ynormal, c="orange", linewidth =1)
```

```
xoverweight=[1.45,2.05] #Min and Max heights in the dataset
yoverweight=[63,126] #calculated coordinates for each cathegory of BMI
figure1= plt.plot(xoverweight, yoverweight, c="green", linewidth =1)
```

```
xobesity2=[1.45,2.05] #Min and Max heights in the dataset
yobesity2=[73.5,147] #calculated coordinates for each cathegory of BMI
figure1= plt.plot(xobesity2, yobesity2, c="orange", linewidth =1)
```

```
xobesity3=[1.45,2.05] #Min and Max heights in the dataset
yobesity3=[84.1,168.1] #calculated coordinates for each cathegory of BMI
figure1= plt.plot(xobesity3, yobesity3, c="red", linewidth =1)
```



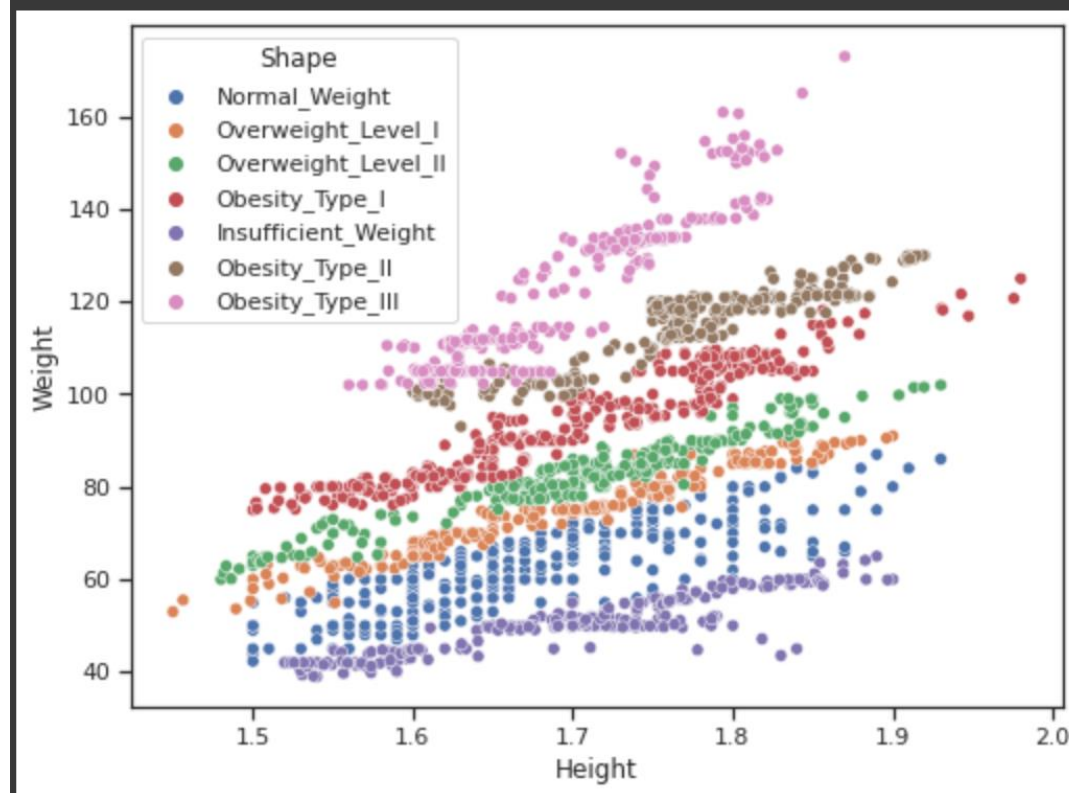


Let's see if this is relevant

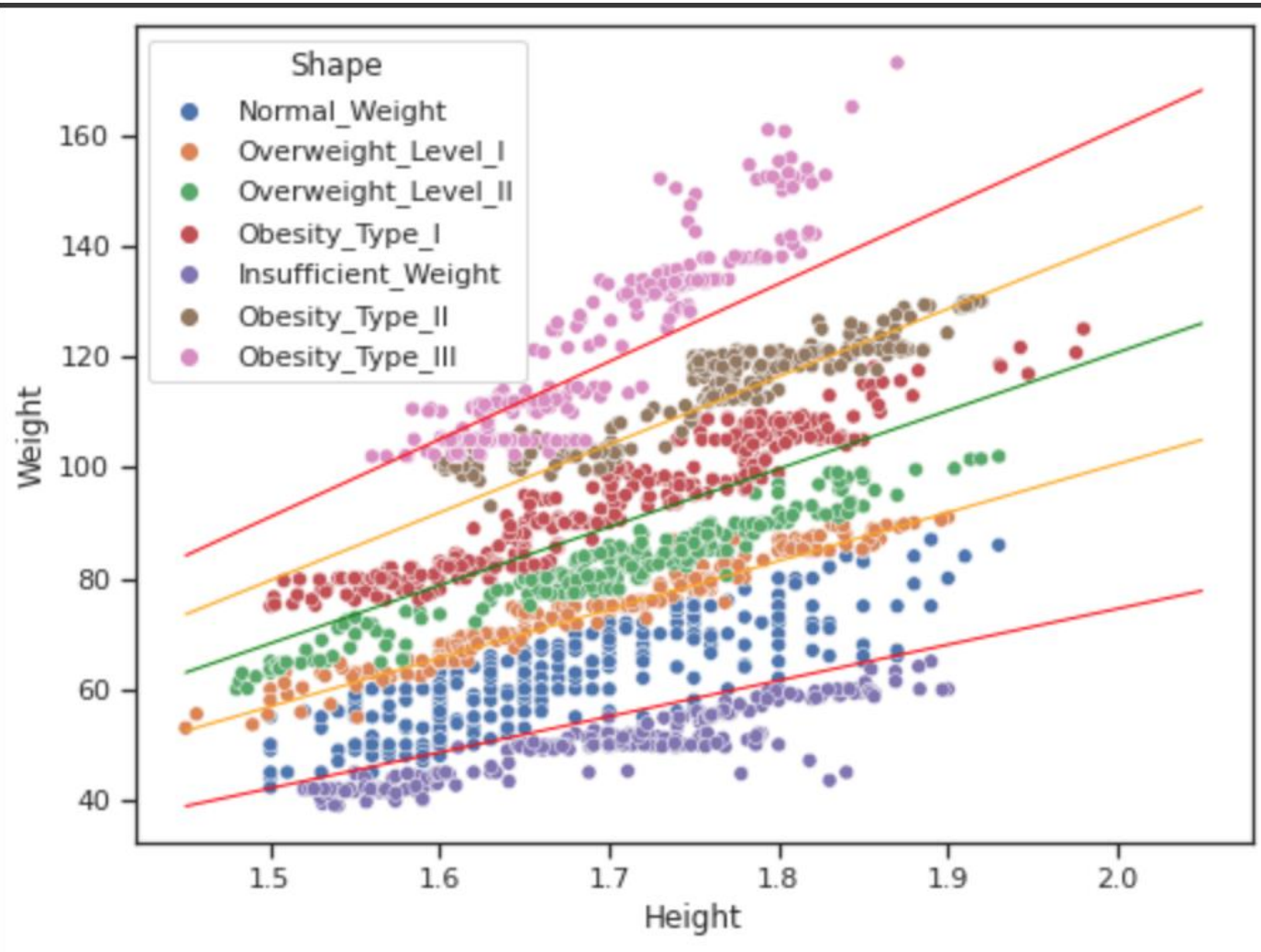
Comparing to the data, BMI sections go the same way as the obesity level classification !

Both graphs looks quite the same.

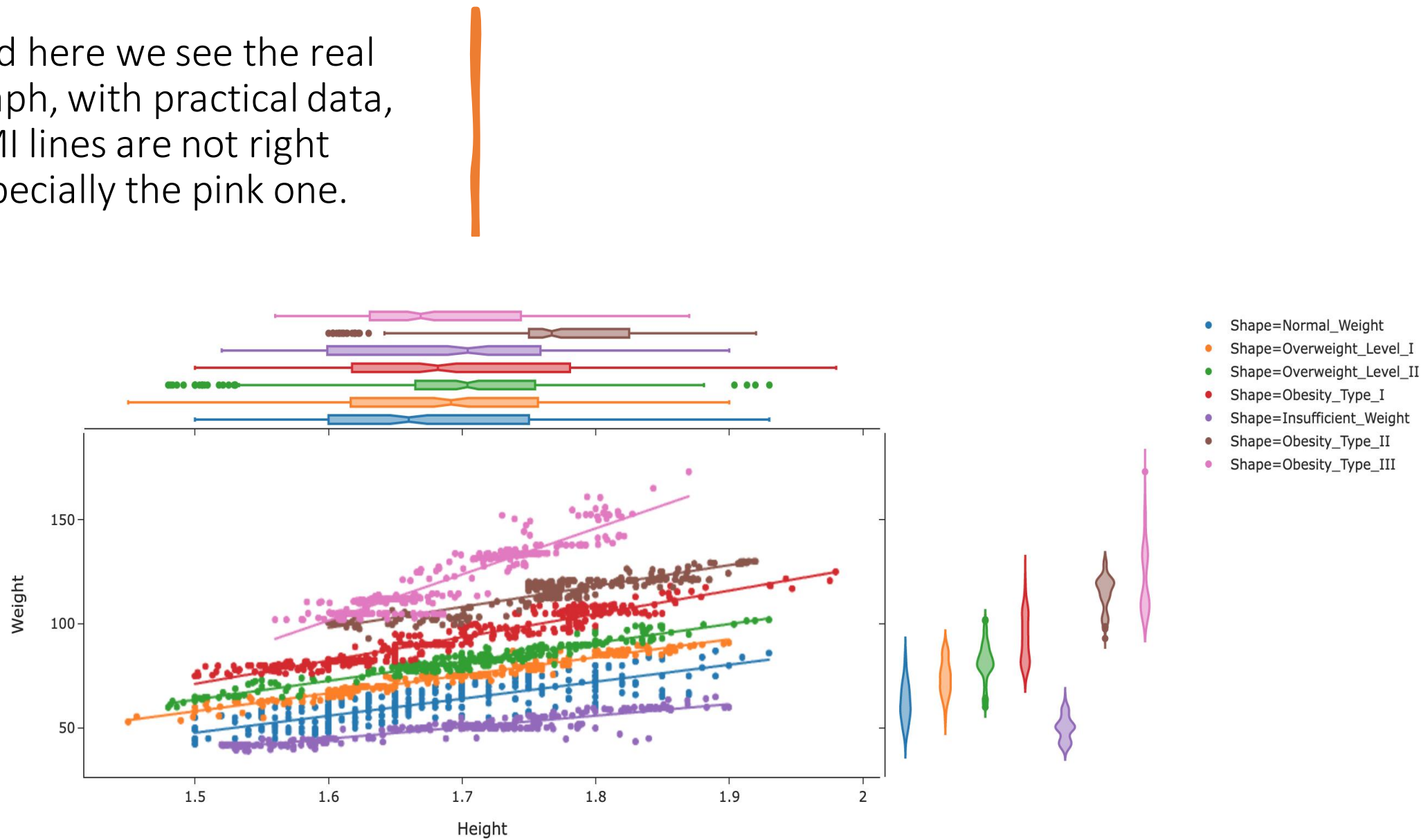
```
f, ax = plt.subplots(figsize=(8, 6))  
  
sns.scatterplot(x='Height', y='Weight',  
               hue='Shape', data=df);
```



Let's
combine
them to see
it clearer !



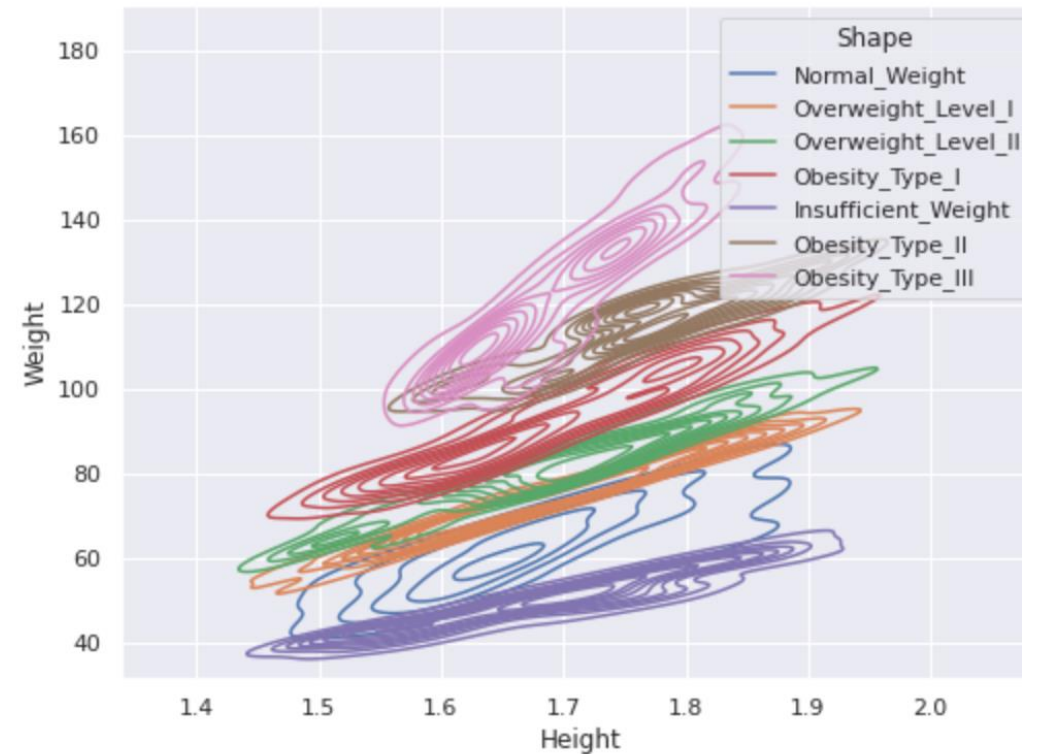
And here we see the real graph, with practical data, BMI lines are not right especially the pink one.



It seems that there is a little problem

- On this plot showing the density of each shape, we can clearly see that they do overlap.
- This is not supposed to happen because the result is determined only with the BMI using weight and height so the limits of each shape should be exact.
- However, the information about the dataset says that the shape is only determined with BMI (height and weight). There is a contradiction.

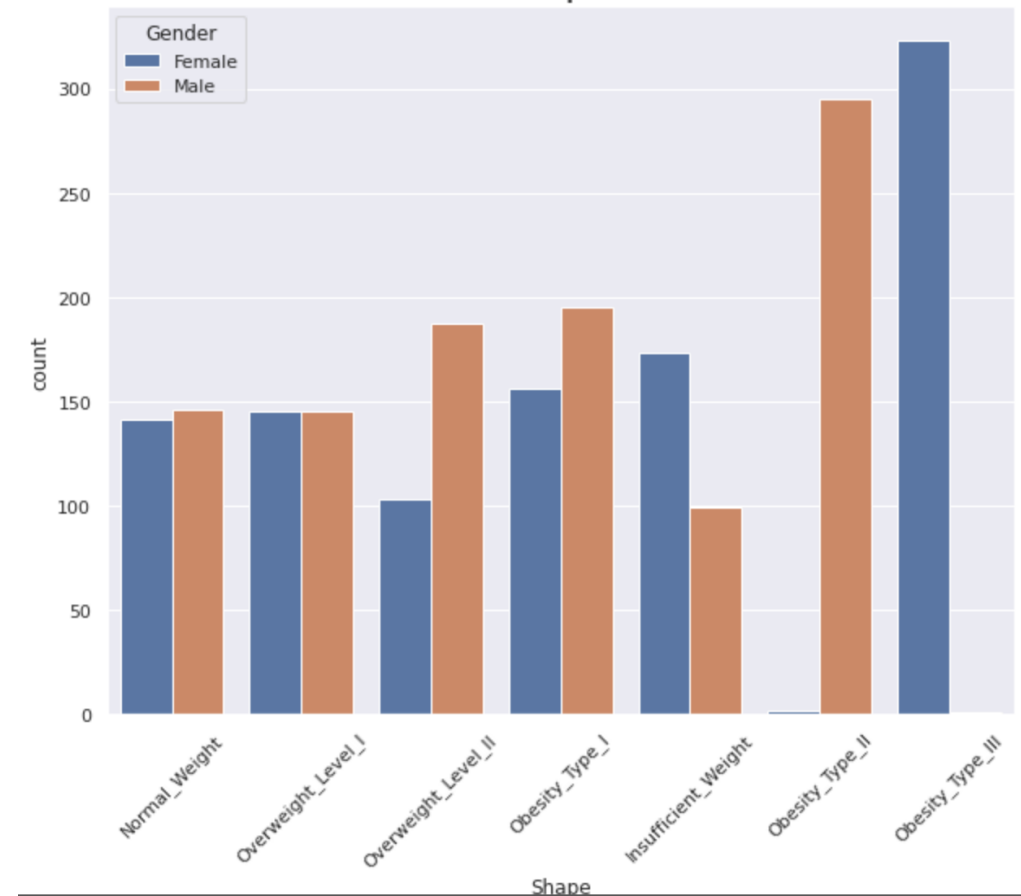
```
# Set up the figure
f, ax = plt.subplots(figsize=(8, 6))
# Draw a contour plot to represent each bivariate density
sns.kdeplot([
    data=df,
    x="Height",
    y="Weight",
    hue="Shape",])
<matplotlib.axes._subplots.AxesSubplot at 0x7fca602fea90>
```



Another anomaly in the dataset

It appears on this graph, we can see that almost only men are affected by obesity level 2 and on the contrary only women are affected by obesity level 3

```
f, ax = plt.subplots(figsize=(10, 8))  
  
sns.countplot(x= 'Shape', hue = 'Gender', data = df)  
plt.title('Shape', weight='bold')  
plt.xticks(rotation=45)
```

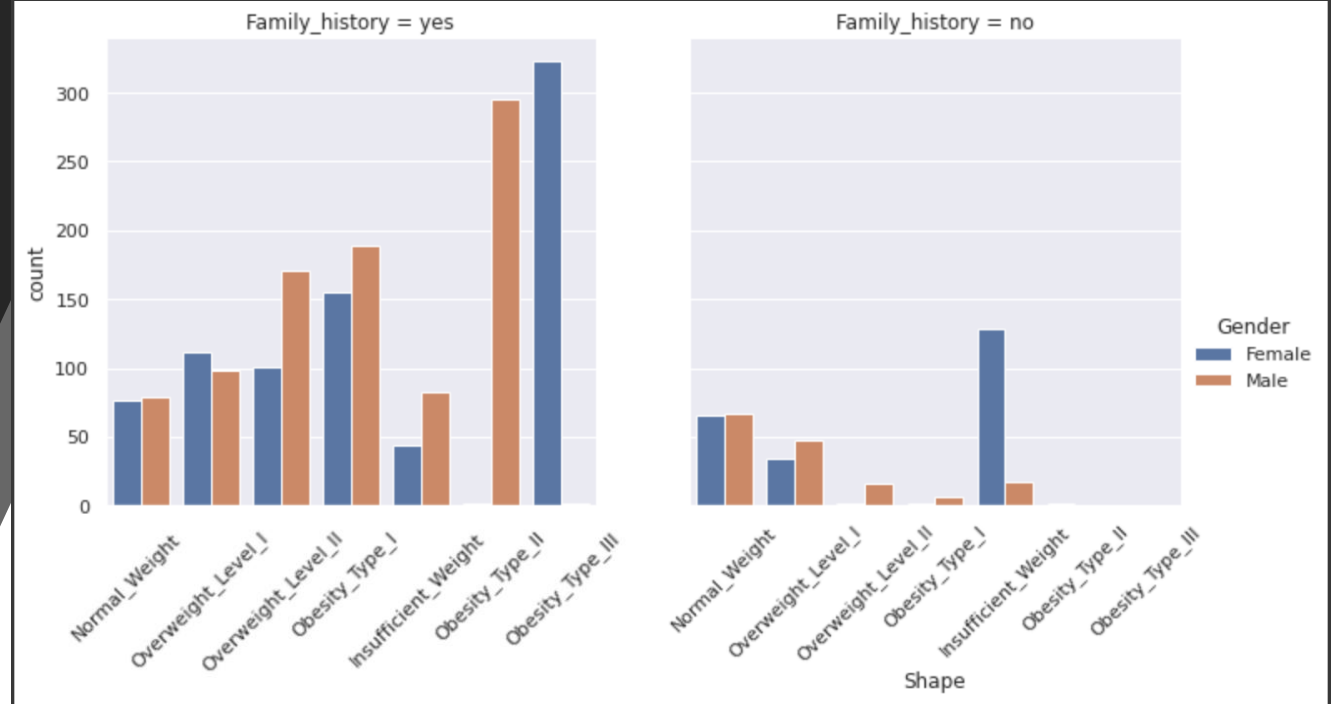


Family history, a major factor in obesity

- We quickly notice that the more obese you are, the more often you have a family history, so much so that only those with a family history were able to have level 2 or 3 obesity in this database.
- However, no family history seems to involve more insufficient weight

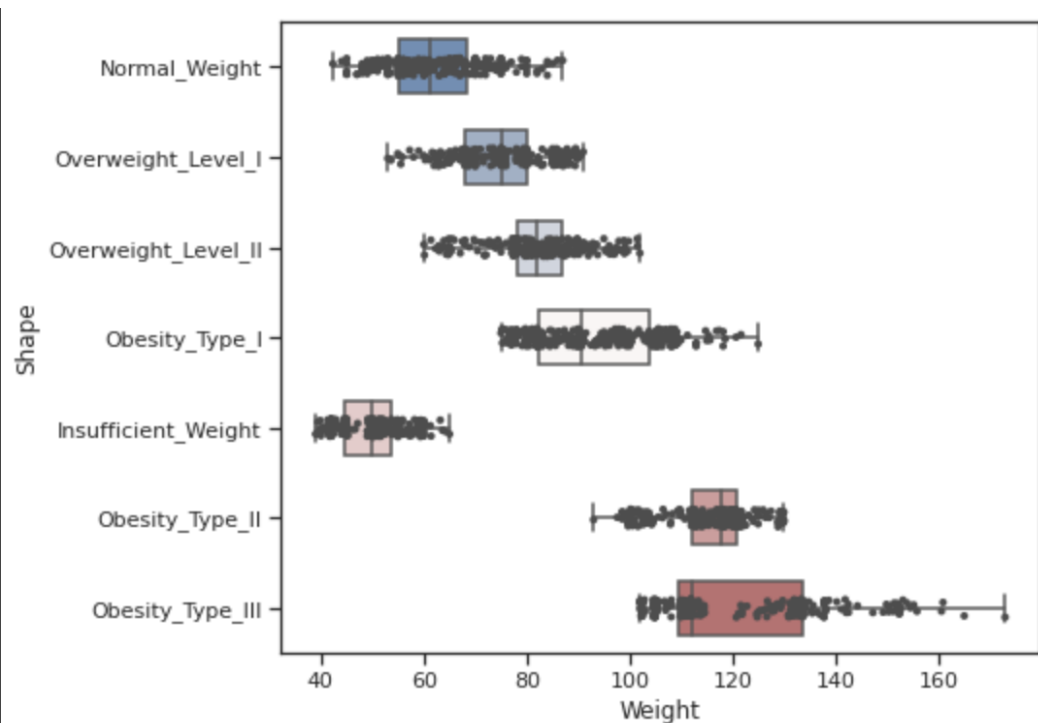
```
sns.catplot(x='Shape', hue="Gender", col="Family_history", data = df, kind="count");  
plt.xticks(rotation=45)
```

```
(array([0, 1, 2, 3, 4, 5, 6]), <a list of 7 Text major ticklabel objects>)
```



BOXPLOTS

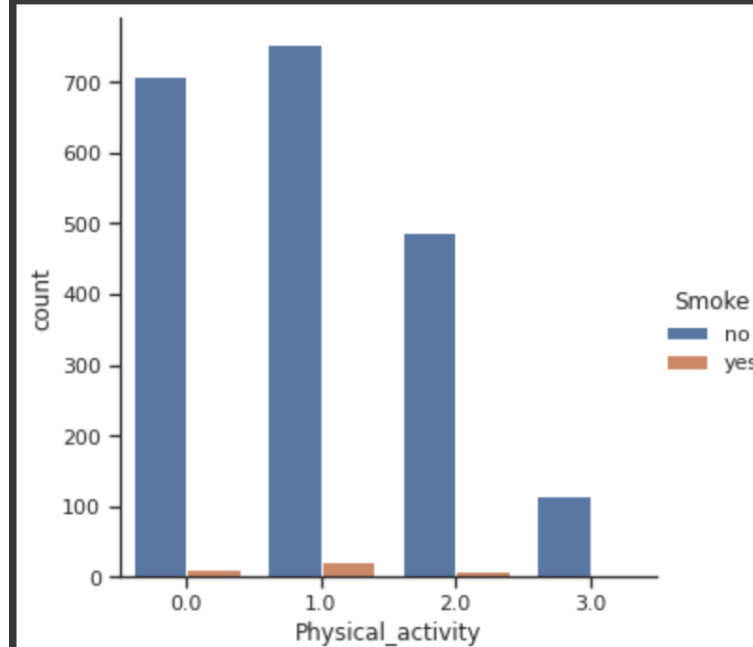
- Boxplots show us that if we get the weight of someone, we can already tell with a high efficiency in which categories can be the person.



```
sns.set_theme(style="ticks")
# Initialize the figure with a logarithmic x axis
f, ax = plt.subplots(figsize=(7, 6))
# Plot the orbital period with horizontal boxes
sns.boxplot(x="Weight", y="Shape", data=df,
            whis=[0, 100], width=.6, palette="vlag")
# Add in points to show each observation
sns.stripplot(x="Weight", y="Shape", data=df,
              size=4, color=".3", linewidth=0)
# Tweak the visual presentation
ax.xaxis.grid(True)
ax.set(ylabel="")
sns.despine(trim=True, left=True)
```

Smoking and physical activity

```
dfsport=df.round({'Physical_activity': 0})  
sns.catplot(x='Physical_activity', hue="Smoke", data = dfsport, kind="count");
```

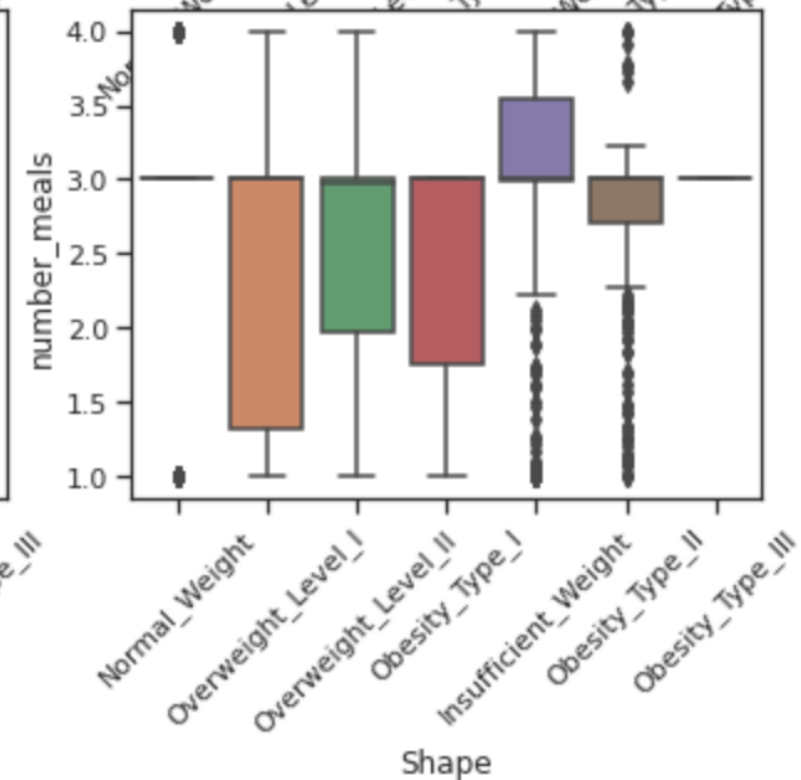
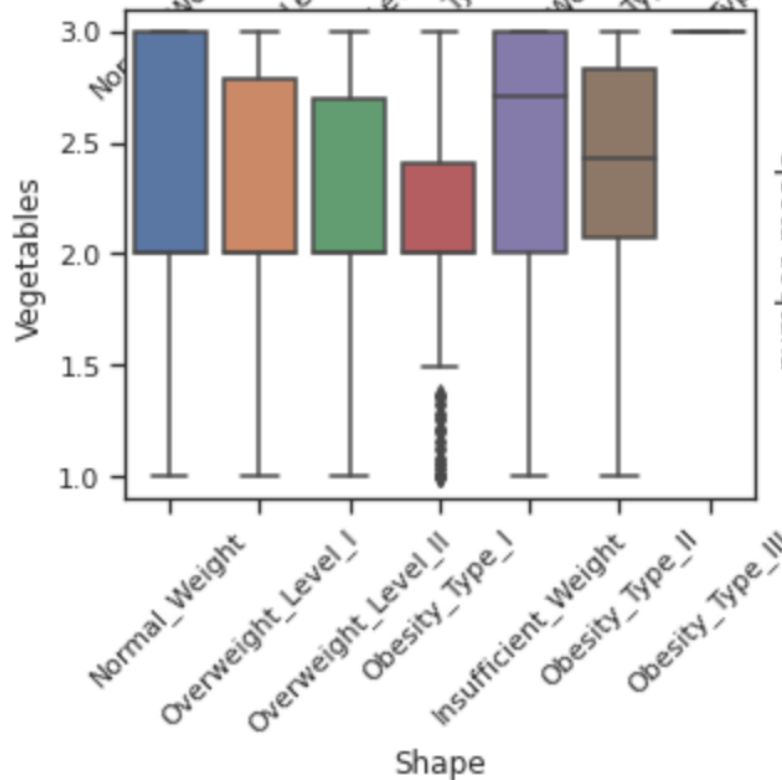
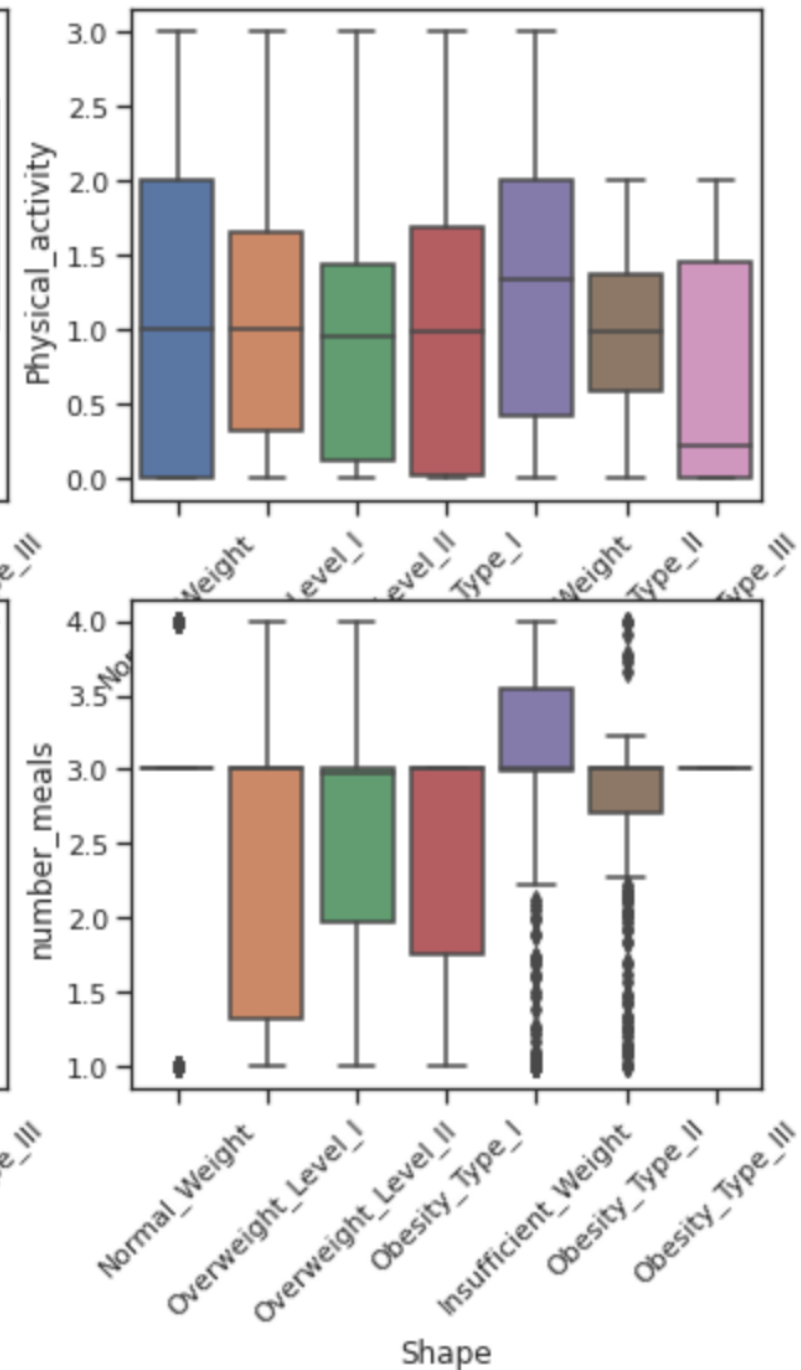
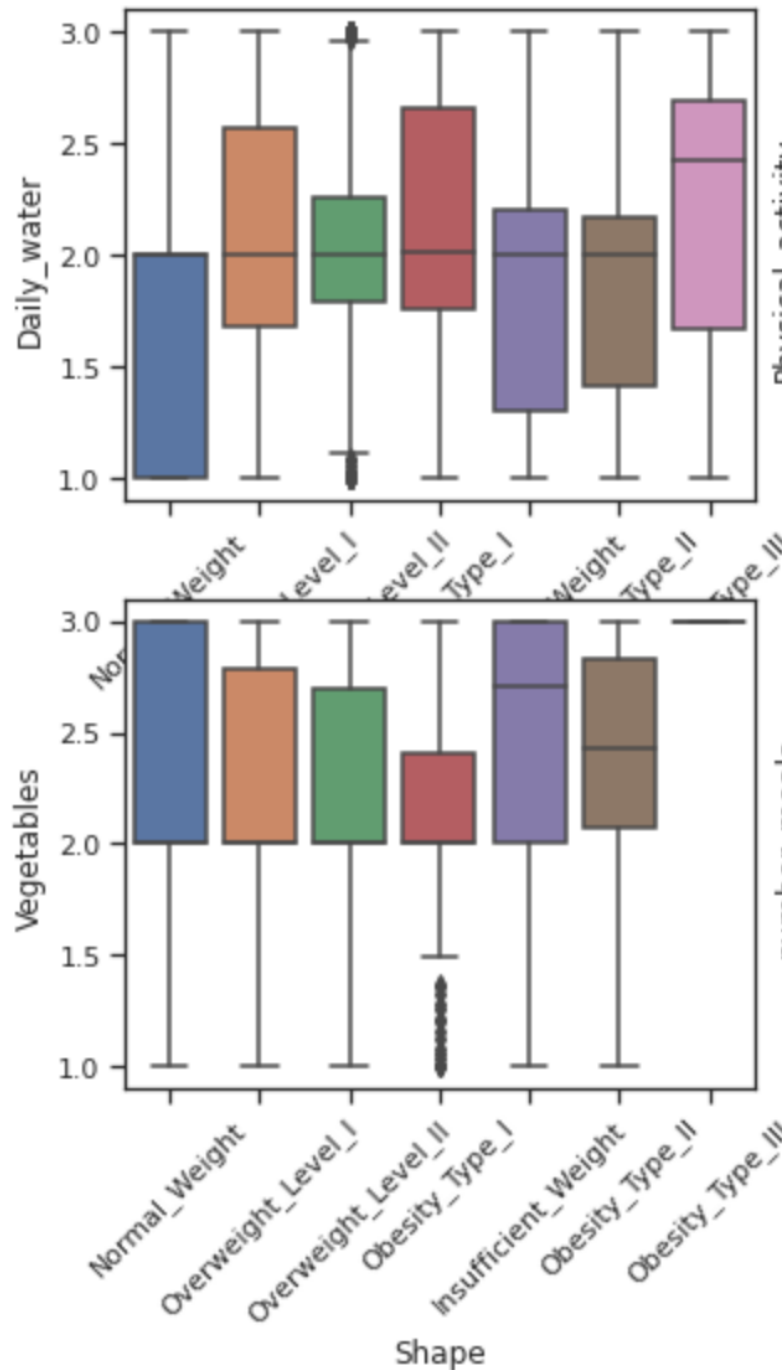


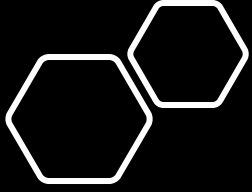
- - People doesn't smoke a lot in this dataset, only 44 smokers out of nearly 2200 !
- - Also, we can see that no smoker play sports more than twice a week.

Comparison of a precise characteristic between different obesity levels

We can see that they are all useful in their way.

```
[115] f, ax = plt.subplots(figsize=(10,8))
plt.subplot(221)
sns.boxplot(x="Shape", y="Daily_water", data=df)
plt.xticks(rotation = 45)
plt.subplot(222)
sns.boxplot(x="Shape", y="Physical_activity", data=df)
plt.xticks(rotation = 45)
plt.subplot(223)
sns.boxplot(x="Shape", y="Vegetables", data=df)
plt.xticks(rotation = 45)
plt.subplot(224)
sns.boxplot(x="Shape", y="number_meals", data=df)
plt.xticks(rotation = 45)
```





2) Data classification

- In this part, we will use different machine learning process.
- First of all, we have to encode to just have numeric values.

K- Neighbors

```
import plotly.express as px
import numpy as np
from sklearn.datasets import make_moons
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier

data=data_dummies.drop(['Age'],axis=1)

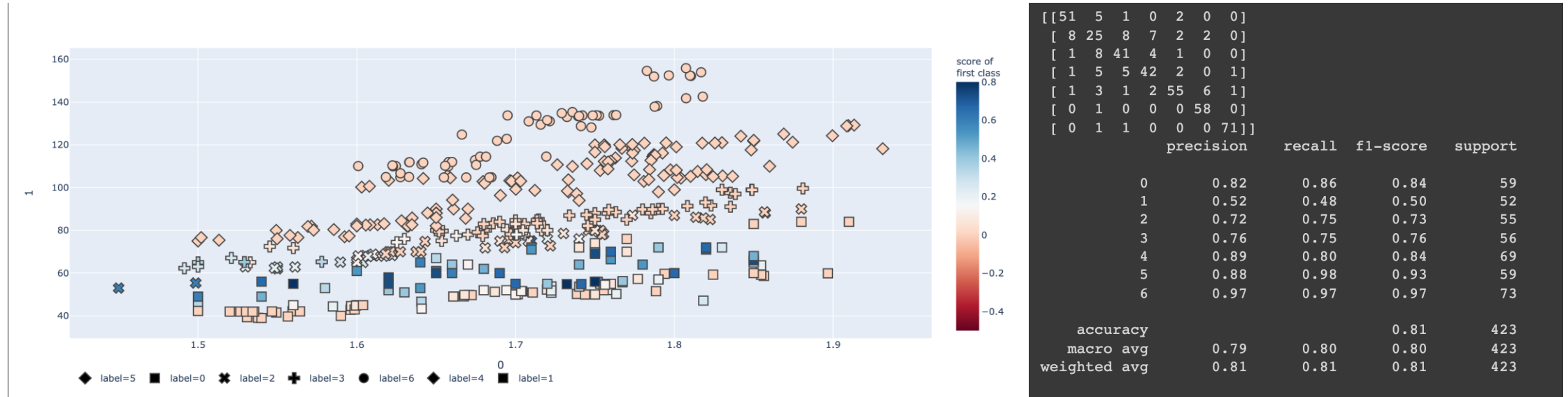
X = data.iloc[:, data.columns != 'Shape'].values
y = data.iloc[:, 7].values

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.20)

# Fit the model on training data, predict on test data
clf = KNeighborsClassifier(15)
clf.fit(X_train, y_train)
y_score = clf.predict_proba(X_test)[: , 1]

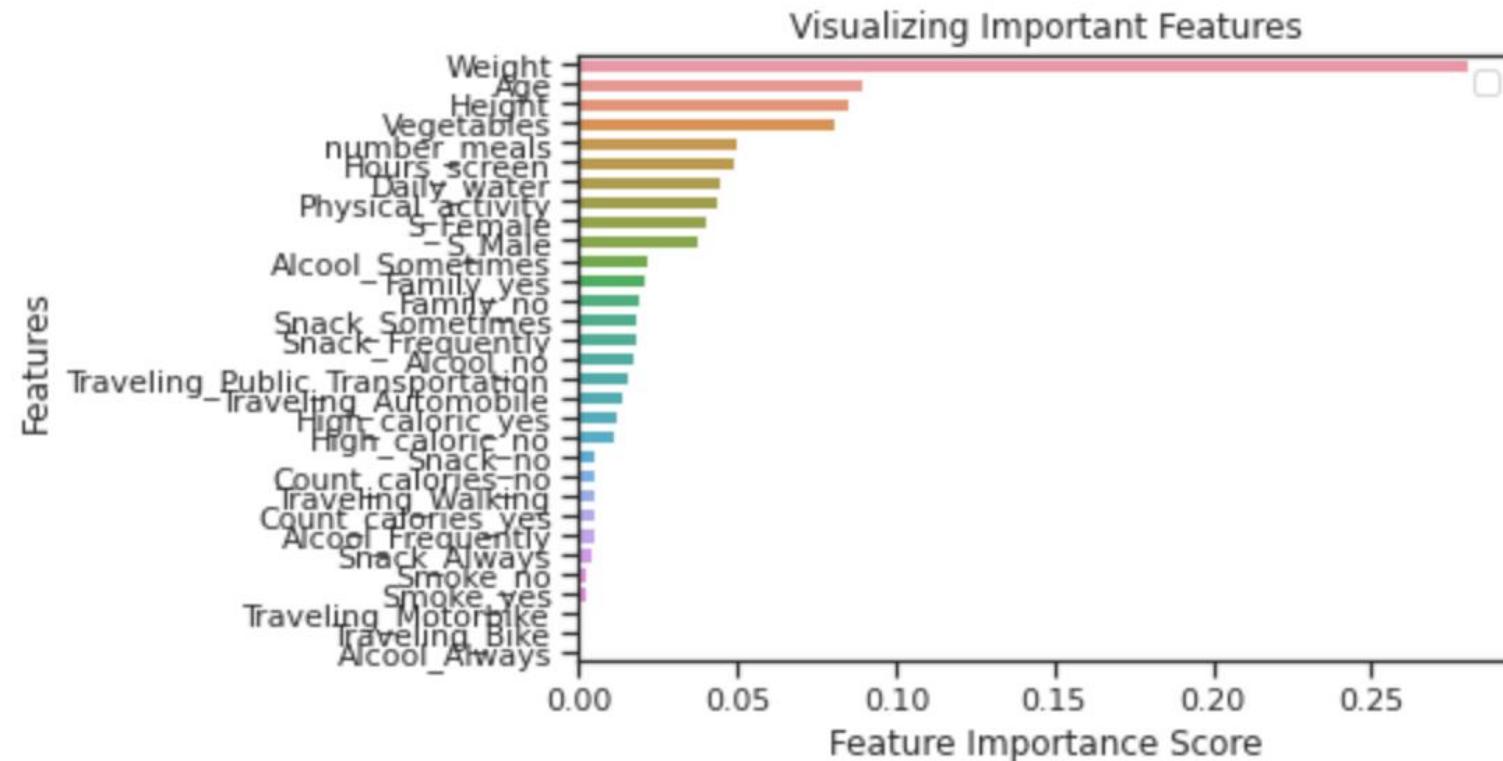
fig = px.scatter(
    X_test, x=0, y=1,
    color=y_score, color_continuous_scale='RdBu',
    symbol=y_test, symbol_map={'0': 'square-dot', '1': 'circle-dot', '2': 'circle',
                              '3': 'square', '4': 'cross-dot', '5': 'cross'},
    labels={'symbol': 'label', 'color': 'score of <br>first class'})
fig.update_traces(marker_size=12, marker_line_width=1.5)
fig.update_layout(legend_orientation='h')
fig.show()
```

K- Neighbors



Here is the visualized results of the classification. It actually worked well the accuracy is good and the representation looks like the one we had at the start.

Random Forest



```
[161] f, ax = plt.subplots(figsize=(10,8))
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
# Creating a bar plot
sns.barplot(x=feature_imp, y=feature_imp.index)
# Add labels to your graph
plt.xlabel('Feature Importance Score')
plt.ylabel('Features')
plt.title("Visualizing Important Features")
plt.legend()
plt.show()
```

We can say with no surprise that the weight is the most important factor of obesity, age height and eating vegetable are also very important.

```
22/22 [=====] - 0s 1ms/step - loss: -523756.2188 - accuracy: 0.1478  
[-523756.21875, 0.14777618646621704]
```

```
Epoch 1/8  
1414/1414 [=====] - 3s 2ms/step - loss: -485.0847 - accuracy: 0.1266  
Epoch 2/8  
1414/1414 [=====] - 2s 2ms/step - loss: -7638.5488 - accuracy: 0.1301  
Epoch 3/8  
1414/1414 [=====] - 2s 2ms/step - loss: -28942.5684 - accuracy: 0.1301  
Epoch 4/8  
1414/1414 [=====] - 2s 2ms/step - loss: -67270.6953 - accuracy: 0.1301  
Epoch 5/8  
1414/1414 [=====] - 2s 2ms/step - loss: -125427.8672 - accuracy: 0.1301  
Epoch 6/8  
1414/1414 [=====] - 2s 2ms/step - loss: -206218.9531 - accuracy: 0.1301  
Epoch 7/8  
1414/1414 [=====] - 2s 2ms/step - loss: -310853.0625 - accuracy: 0.1301  
Epoch 8/8  
1414/1414 [=====] - 2s 2ms/step - loss: -443123.0938 - accuracy: 0.1301  
<keras.callbacks.History at 0x7f52c86152d0>
```

Neural Network

The neural network was very hard to set up. The results obtained are very bad. We see an accuracy of 14% which means that the code does not work. We would have liked with more time to understand the reason for this failure.



Thank you for your
attention