Our ultimate goal is to minimize the training error.

Another way to view this observation is that our ultimate goal, is to extract a hypothesis, which will return the minimum training error.

# **Perceptron Algorithm**

In this section, we are going to introduce the percepton algorithm. Perceptron algorithm, will generate a hypothesis function (linear classifier) for a given data set under consideration. From the explanation of the perceptron algorithm, we may receive a wealth of introductory needed concepts.

# **Introductory Example**

Task: Given 2-d observation data points, return a hypothesis (using perceptron algorithm) that classifies them in the best possible way

```
In [3]: # Package for scientific computing with Python
# https://numpy.org/
import numpy as np

# Matplotlib is a library for creating static, animated, and interactive visualizations in Python.
# https://matplotlib.org/
import matplotlib.pyplot as plt

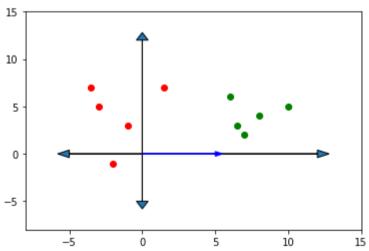
# our horizontal and vertical graph axes
ax = plt.axes()

# x - axis positive:
# origin_vector_x, origin_vector_y, dx, dy
ax.arrow(0.0, 0.0, 12, 0.0, head_width=0.8, head_length=0.8)

# x - axis negative:
# origin_vector_x, origin_vector_y, dx, dy
ax.arrow(0.0, 0.0, -5.0, 0.0, head_width=0.8, head_length=0.8)

# y-axis positive
```

```
ax.arrow(0.0, 0.0, 0.0, 12.0, head_width=0.8, head_length=0.8)
#ax.text(-7, 4, "theta vector")
# y-axis negative
ax.arrow(0.0, 0.0, 0.0, -5.0, head width=0.8, head length=0.8)
x_no = np.array([-1, -2, -3, -3.5, 1.5])
y_{no} = np.array([3, -1, 5, 7, 7])
x \text{ yes} = [7, 6, 6.5, 8, 10]
y_yes = [2, 6, 3, 4, 5]
# y axes of figure limits
plt.ylim(-8,15)
# x axes of figure limits
plt.xlim(-8,15)
plt.plot(x no,y no,'o', color='red')
plt.plot(x yes,y yes,'o', color='green')
ax.arrow(0.0,0.0, 5.0, 0.0, head width=0.5, head length=0.5, color="blue")
plt.show()
```



The horizontal axis (x) contains  $x_1^{(i)}$  and the vertical axis (y) contains the  $x_2^{(i)}$ . With red color are depicted the data points labeled as -1, whereas with green color are depticted the data points labeled as +1.

### Initialization

Based uppon the concepts of the previous introductory section, we are going to arbitrarily initialize a  $\theta$  vector as well as a  $\theta_0$  quantity as follows:

$$\bar{\theta} = (5.0, 0.0),$$
 $\theta_0 = 0.0$ 

Additionally, we are going to initialize a constant value m, which will define our maximum number of iterations.

```
In [4]: # maximum number of iterations (a.k.a. epochs)
m = 10
```

### Data Set

In order for our algorithm to function, we have to pass a labeled data set. Therefore, we will give for each observation data point a label and then we are going to merge all data points into a general set.

```
In [5]: # red points labeled as -1
    x_no = np.array([-1, -2, -3, -3.5, 1.5])
    y_no = np.array([-1, -1, -1, -1, -1])
    label_no = np.array([-1, -1, -1, -1, -1])

# green points labeled as +1
    x_yes = [7, 6, 6.5, 8, 10]
    y_yes = [2, 6, 3, 4, 5]
    label_yes = np.array([1, 1, 1, 1])

In [6]: # merge x-coordinates of all observations into 1 set
    x = np.append(x_no, x_yes)
    print(x)

[-1, -2, -3, -3.5, 1.5, 7, 6, 6.5, 8, 10, ]

In [7]: # merge y-coordinates of all observations into 1 set
    y = np.append(y_no, y_yes)
    print(y)
```

```
In [8]: # merge all known facts (all labels) into 1 set
    labels = np.append(label_no, label_yes)
    print(labels)

[-1 -1 -1 -1 1 1 1 1 1]

In [9]: # formally initialize theta vector and theta_0 vector
    theta = [5.0, 0.0]
    theta_0 = 0.0
```

# **Perceptron Algorithm**

What follows, is the perceptron algorithm (in pseudo code), the explanation and some insight analysis of its functionality as well as a python - code representation alongside the intermediate steps display of the functionality.

#### Pseudo - Code

```
\begin{array}{c} \text{if(theta\_change == False)} \\ \text{break} \\ \text{return } \overset{-}{\theta}, \theta_0 \end{array}
```

## **Analysis**

There are 2 primary points of interest worth noting in our perceptron algorithm:

1. The first is, when does our perceptron algorithm changes or is being modified.

We can see that perceptron algorithm is being modified whenever this condition becomes true

$$y^{(i)} \cdot (\overline{ heta^T} \cdot \overline{x^{(i)}} + heta_0) \leq 0$$

We remember that  $y^{(i)}$  is our label (or established fact), and can receive either a positive number +1, or a negative number -1. Furthemore, we remember that  $\overline{\theta^T} \cdot \overline{x^{(i)}} + \theta_0$  is our prediction hypothesis for a given observation data point  $\overline{x^{(i)}}$ .

Given that our prediction follows the same logic as our labels, classifying either +1 or -1, the prementioned condition becomes true iff: our prediction and the established known fact do not match with each other.

1. The second point of interest worth noting, is how the perceptron is being modified when hypothesis result and label do not match with each other. We see that 2 main modifications take place:

$$\overline{ heta}_{new} = \overline{ heta}_{old} + (y^{(i)} \cdot \overline{x^{(i)}})$$

as well as

$$heta_{0_{new}} = heta_{0_{old}} + y^{(i)}$$

Let us review how the condition of (1) is being modified with the updated values  $\theta_{new}$  and  $\theta_{0_{new}}$ . The condition now becomes:

$$y^{(i)}\cdot(\overline{ heta_{new}^T}\cdot\overline{x^{(i)}}+ heta_{0_{new}})\Rightarrow \ y^{(i)}\cdot[\overline{[ heta_{old}+(y^{(i)}\cdot\overline{x^{(i)}})]^T}\cdot\overline{x^{(i)}}+( heta_{0_{old}}+y^{(i)})]\Rightarrow \ y^{(i)}\cdot[(\overline{ heta_{old}^T}\cdot\overline{x^{(i)}})+(y^{(i)}\cdot\overline{x^{(i)^T}}\cdot\overline{x^{(i)}})+ heta_{0_{old}}+y^{(i)}]\Rightarrow \ y^{(i)}\cdot\overline{ heta_{old}^T}\cdot\overline{x^{(i)}}+y^{(i)^2}\cdot\overline{x^{(i)^T}}\cdot\overline{x^{(i)}}+y^{(i)}\cdot heta_{0_{old}}+y^{(i)^2}\Rightarrow \ y^{(i)}\cdot(\overline{ heta_{old}^T}\cdot\overline{x^{(i)}}+ heta_{0_{old}})+y^{(i)^2}\cdot(\overline{x^{(i)^T}}\cdot\overline{x^{(i)}}+1)\Rightarrow \ y^{(i)}\cdot hypothesis_{old}(\overline{x^{(i)}})+(\|x^{(i)}\|^2+1)$$

What all the prementioned statements actually indicate, is that for every observation data point  $(x^{(i)})$  that has been wrongfully classified ( $y^{(i)} \cdot hypothesis_{old}(\overline{x^{(i)}}) \leq 0$ ), we add a strictly positive quantity based uppon  $x^{(i)}$ , to minimize the distance of  $y^{(i)} \cdot hypothesis_{old}(\overline{x^{(i)}})$  from 0, untill it becomes a positive value  $\forall \overline{x^{(i)}}$ 

Next we are going to present the execution of perceptron algorithm (in python) presented in the previously displayed data - set.

### **Python Code**

First we are going to define a function, **plotPerceptronResult()**, which will generate the graphical representation of the  $\theta_{vector}$ , the perpendicular to  $\overline{\theta}$  vector ( $hypothesis(\overline{v}) = 0$ ) alongside all of our observation data points.

```
In [10]: def plotPerceptronResults(theta, x_i_1, x_i_2):
    # 1. Our horizontal and vertical graph axes
    ax = plt.axes()

# 2. Cartesian axes vectors
    # x - axis positive:
    # origin_vector_x, origin_vector_y, dx, dy
ax.arrow(0.0, 0.0, 12, 0.0, head_width=0.8, head_length=0.8)
    # x - axis negative:
ax.arrow(0.0, 0.0, -5.0, 0.0, head_width=0.8, head_length=0.8)
    # y-axis positive
```

```
ax.arrow(0.0, 0.0, 0.0, 12.0, head width=0.8, head length=0.8)
    # y-axis negative
ax.arrow(0.0, 0.0, 0.0, -5.0, head_width=0.8, head_length=0.8)
# 3. Data
    # -1 labeled red point observations
x \text{ no = np.array}([-1, -2, -3, -3.5, 1.5])
y no = np.array([3, -1, 5, 7, 7])
    # +1 labeled red point observations
x \text{ yes} = [7, 6, 6.5, 8, 10]
y \text{ yes} = [2, 6, 3, 4, 5]
    # data plot
plt.plot(x no,y no,'o', color='red')
plt.plot(x yes,y yes,'o', color='green')
# 4. Figure Limits
    # y axes limits of figure
plt.ylim(-8,15)
    # x axes of figure limits
plt.xlim(-8,15)
# 5. Vectors
    # theta vector
ax.arrow(0.0,0.0, theta[0], theta[1], head width=0.5, head length=0.5, color="blue")
    # perpendicular vector [x1, x2] with x1=5
    # for theta = [c1, c2]^T
    \# c1*x1 + c2*x2 = 0 = >
    \# c1*5 + c2*x2 = 0 = >
    \# c2*x2 = -c1*5
    \# x2 = (-c1/c2) * 5
    # plot from (0,0) to (5, (-c1/c2) * 5)
ax.arrow(0.0,0.0, 5.0, (-theta[0]/theta[1])*5, head_width=0.5, head_length=0.5, color="red")
    #symmetric
    # plot from (0,0) to (-5, (c1/c2) * 5)
ax.arrow(0.0,0.0, -5.0, (theta[0]/theta[1])*5, head width=0.5, head length=0.5, color="red")
# 6. Point under consideration
plt.plot(x_i_1, x_i_2, 'v', color="black")
plt.show()
```

Next, we are defining a function, **dotProduct()**, which performs vector multiplication between a given  $\theta$  and a given  $x^{(i)}$ , where  $\theta, x^{(i)} \in \mathbb{R}^2$ 

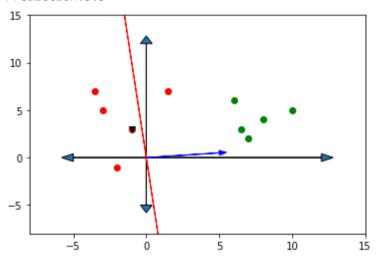
```
In [11]: # function that receives two vectors that can
    # be multiplied and returns their dot product

def dotProduct(theta_vector, data_point_x, data_point_y):
    result = 0
    result += theta_vector[0]*data_point_x
    result += theta_vector[1]*data_point_y
    return result
```

```
Finally we present, the perceptron algorithm:
In [12]: # formally initialize theta vector and theta 0 vector
          theta = [5.0, 0.5]
          theta 0 = 0.0
In [13]: # iterate from 1 to m
         for t in range(1, m+1):
              print("Iteration:"+str(t))
              # have we observed a change in theta vector
              is theta changed = False
              #iterate through data points
              for i in range(len(x)):
                  prediction = labels[i] * dotProduct(theta, x[i], y[i]) + labels[i] * theta 0
                  print("Observation data point:["+str(x[i])+","+str(y[i])+"]")
                  print("Label"+str(labels[i]))
                  print("Theta:["+str(theta[0])+","+str(theta[1])+"]")
                  print("Theta 0:"+str(theta 0))
                  print("Prediction:"+str(prediction))
                  plotPerceptronResults(theta, x[i], y[i])
                  if(prediction<=0):</pre>
                      #our hypothesis prediction is wrong and has to change
                      print("Wrong prediction")
                      theta[0] += labels[i]*x[i]
                      theta[1] += labels[i]*y[i]
                      theta_0 += labels[i]
```

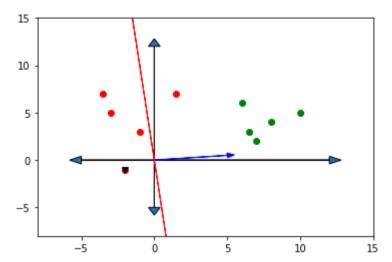
```
print("New Theta:["+str(theta[0])+","+str(theta[1])+"]")
           print("New theta_0:"+str(theta_0))
           is_theta_changed = True
       else:
           print("Right prediction")
       print("----\n")
    if(is theta changed==False):
       break
    print("======")
Iteration:1
Observation data point:[-1.0,3]
Label-1
```

Theta:[5.0,0.5] Theta 0:0.0 Prediction:3.5



Right prediction

Observation data point:[-2.0,-1] Label-1 Theta:[5.0,0.5] Theta\_0:0.0 Prediction:10.5



## Right prediction

-----

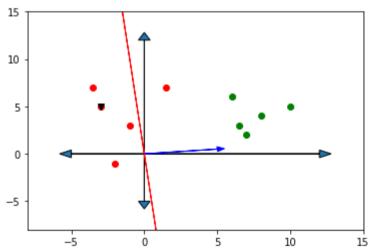
Observation data point:[-3.0,5]

Label-1

Theta:[5.0,0.5]

Theta\_0:0.0

Prediction:12.5



# Right prediction

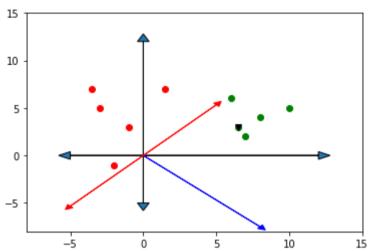
Observation data point:[6.5,3]

Label1

Theta:[8.0,-7.5]

Theta\_0:-1.0

Prediction:28.5



Right prediction

-----

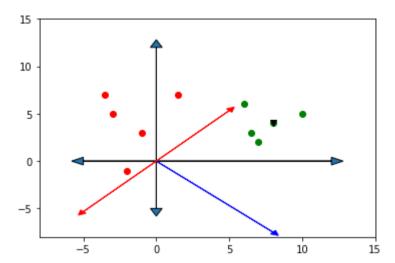
Observation data point:[8.0,4]

Label1

Theta:[8.0,-7.5]

Theta\_0:-1.0

Prediction:33.0



## Right prediction

-----

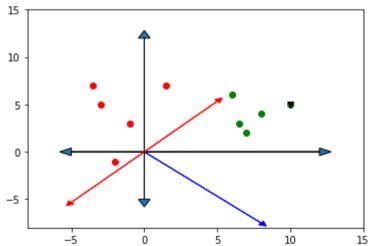
Observation data point:[10.0,5]

Label1

Theta:[8.0,-7.5]

Theta\_0:-1.0

Prediction:41.5



Right prediction

-----

It is easy to observe, that if our data set of observations is linearly classified through the origin, then the perceptron algorithm can be executed as it is, with no adjustments or modifications. Nevertheless, we can be certain that it is extremely likely to face an example of a data - set of observations, that is linearly classifiable, nevertheless it is not linearly classifiable through the origin. Let us review an alike example as well as the proper modifications that we have to make.

# **Basketball Example**

Let us assume, that we are given as a task to generate a hypothesis based uppon the following:

#### Task:

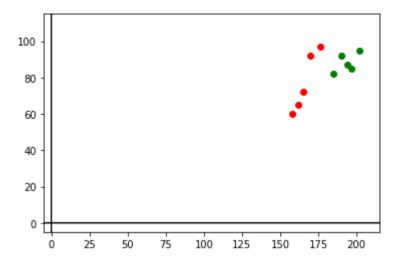
We want to extract a prediction on whether or not, a high - scholl student of a given high - school, will be picked/drafted to take part in the High - School Basketball team for the current championship season, based uppon students height (in cms) and weight (in kg). We are given an observation data set, of (height, weight) vectors of students that were successfully picked as well as not picked, during the last 3 years.

#### Data Set:

#Student Observation $\boldsymbol{x}^{(i)}$	Height (cm-s)	Weight (kg-s)	Chosen (label)
$x^{(0)}$	158	60	-1
$x^{(1)}$	165	72	-1
$x^{(2)}$	170	92	-1
$x^{(3)}$	162	65	-1
$x^{(4)}$	176	97	-1
$x^{(5)}$	185	82	+1
$x^{(6)}$	194	87	+1
$x^{(7)}$	190	92	+1
$x^{(8)}$	197	85	+1
$x^{(9)}$	202	95	+1

#### Data Plot:

```
In [14]: # 1. Our horizontal and vertical graph axes
         ax = plt.axes()
         # 2. Cartesian axes vectors
             # x - axis positive:
             # origin vector x, origin vector y, dx, dy
         ax.arrow(0, 0, 215.0, 0.0, head width=0.8, head length=0.8)
             # x - axis negative:
         ax.arrow(0.0, 0.0, -5.0, 0.0, head width=0.8, head length=0.8)
              # y-axis positive
         ax.arrow(0.0, 0.0, 0.0, 115.0, head width=0.8, head length=0.8)
              # y-axis negative
         ax.arrow(0.0, 0.0, 0.0, -5.0, head width=0.8, head length=0.8)
         # 3. Data
             # -1 labeled red point observations
         x \text{ no} = np.array([158, 165, 170, 162, 176])
         y_no = np.array([60, 72, 92, 65, 97])
             # +1 labeled red point observations
         x \text{ yes} = [185, 194, 190, 197, 202]
         y_yes = [82, 87, 92, 85, 95]
             # data plot
         plt.plot(x no,y no,'o', color='red')
         plt.plot(x yes,y yes,'o', color='green')
         # 4. Figure limits
              # y axes limits of figure
         plt.ylim(-5,115)
             # x axes of figure limits
         plt.xlim(-5,215)
         plt.show()
```



We can see by the prementioned plot, that our data set is linearly classified, nevertheless is not linearly classified through the origin.

# Linear classification through the origin

We can transform the existing data - set to be linearly classifiable through the origin, by performing a very simple dimensionanility transformation. More specifically, for:

$$\overset{-}{ heta}=( heta_1, heta_2)$$

and

$$\overline{x^{(i)}} = (x_1^{(i)}, x_2^{(i)})$$

we can increase the dimensions by +1 as follows:

$$rac{ar{ heta}=( heta_1, heta_2, heta_0)}{x^{(i)}=(x_1^{(i)},x_2^{(i)},1)}$$

In order for our human perception to understand more easily the way the algorithm functions via this transformation (also known as bias insertion) we can visualize the data points in  $R^2$ , where we have to expect that  $\forall x^{(i)}$ , the data points will be successfully classified via the equation:

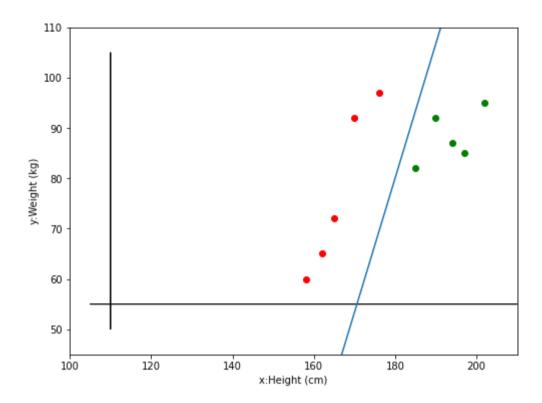
$$heta_1 \cdot x_1^{(i)} + heta_2 \cdot x_2^{(i)} + heta_0 \cdot 1 = 0$$

Let us review the given data set under this point of view, given a  $\theta$  that correctly classifies the observation data set:

$$\bar{\theta} = (40, -15, -6000)$$

```
In [15]: # Package for scientific computing with Python
         # https://numpy.org/
         import numpy as np
         # Matplotlib is a library for creating static, animated, and interactive visualizations in Python.
         # https://matplotlib.org/
         import matplotlib.pyplot as plt
         from mpl toolkits.mplot3d.art3d import Poly3DCollection
In [16]: # -1 labeled red point observations
         x \text{ no} = \text{np.array}([158, 165, 170, 162, 176])
         y \text{ no} = np.array([60, 72, 92, 65, 97])
         z no = np.array([1, 1, 1, 1, 1])
             # +1 labeled red point observations
         x \text{ yes} = [185, 194, 190, 197, 202]
         y \text{ yes} = [82, 87, 92, 85, 95]
         z \text{ yes} = [1, 1, 1, 1, 1]
         theta = [40, -15, -6000]
         fig = plt.figure(figsize=(8,6))
         # 1. Add subplot (rows, columns, index)
         ax = fig.add subplot(111)
         ax.set xlim([100,210])
         # height
         ax.set ylim([45,110])
         # generate x axis
             #positive
         ax.arrow(110, 55, 100, 0.0, color="black")
             #negative
         ax.arrow(110, 55, -5, 0.0, color="black")
```

```
# generate y axis
   #positive
ax.arrow(110, 55, 0.0, 50, color="black")
   #negative
ax.arrow(110, 55, 0.0, -5, color="black")
# 2. Plot all the observation data points
ax.plot(x no, y no, 'o', color='red')
ax.plot(x yes, y yes, 'o', color='green')
ax.set xlabel("x:Height (cm)")
ax.set ylabel("y:Weight (kg)")
#======= GENERATE 2 POINTS FOR THE CLASSIFIER LINE ==========
# equation:
\# x2 = -(x1*theta1 + 1*theta0) / theta2
\# y = -(x1*theta[0] + 1*theta[2]) / theta[1]
# for x = [120, 210]
x = [-120, 210]
y = [0,0]
y[0] = -(x[0]*theta[0] + 1*theta[2]) / theta[1]
y[1] = -(x[1]*theta[0] + 1*theta[2]) / theta[1]
ax.plot(x,y)
plt.show()
```



It would be interesting to review the verification of the correct theta of our perceptron algorithm, by reviewing its 3D representation alongside.

$$\bar{ heta} = (40, -15, -6000)$$

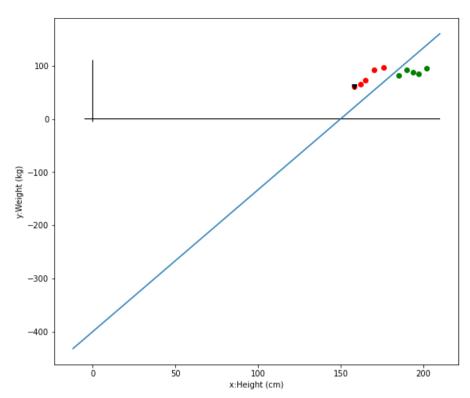
```
#ax.set xlim([100,210])
# heiaht
#ax.set ylim([45,110])
# generate x axis
   #positive
ax.arrow(0, 0, 210, 0.0, color="black")
   #negative
ax.arrow(0, 0, -5, 0.0, color="black")
# generate v axis
   #positive
ax.arrow(0, 0, 0.0, 110, color="black")
   #negative
ax.arrow(0, 0, 0.0, -5, color="black")
# 2. Plot all the observation data points
ax.plot(x no, y no, 'o', color='red')
ax.plot(x yes, v yes, 'o', color='green')
ax.plot(x_i_x, x_i_y, 'v', color='black')
ax.set xlabel("x:Height (cm)")
ax.set ylabel("y:Weight (kg)")
#======= GENERATE 2 POINTS FOR THE CLASSIFIER LINE ===========
# equation:
\# x2 = -(x1*theta1 + 1*theta0) / theta2
\# y = -(x1*theta[0] + 1*theta[2]) / theta[1]
# for x = [120, 210]
x = [-12,210]
y = [0,0]
y[0] = -(x[0]*theta[0] + 1*theta[2]) / theta[1]
y[1] = -(x[1]*theta[0] + 1*theta[2]) / theta[1]
ax.plot(x,y)
# add subplot (3d projection of rows, columns, index)
ax = fig.add subplot(122, projection='3d')
#ax.set_xlim([100,210])
#ax.set_ylim([45,110])
ax.set_zlim([-0.5,1.2])
# generate x axis
   #positive
```

```
ax.quiver(0.0, 0.0, 0.0, 220.0, 0.0, 0.0, arrow length ratio=0.001, alpha=0.5, color="black")
    #negative
ax.quiver(0.0, 0.0, 0.0, -220.0, 0.0, 0.0, arrow length ratio=0.001, alpha=0.5, color="black")
# generate x plane
\#x \ x=[-250,250, 250, -250]
\#x_y=[0, 0, 0, 0]
\#x \ z = [-5.0, -5.0, 5.0, 5.0]
\#vertices = [list(zip(x x, x y, x z))]
#poly = Poly3DCollection(vertices, alpha=0.1, color="black")
#ax.add collection3d(poly)
#-----
# generate v axis
    #positive
ax.quiver(0.0, 0.0, 0.0, 120.0, 0.0, arrow length ratio=0.001, alpha=0.5, color="black")
    #negative
ax.quiver(0.0, 0.0, 0.0, -120.0, 0.0, arrow length ratio=0.001, alpha=0.5, color="black")
# generate y plane
y x=[-250,250, 250, -250]
y y = [-130, -130, 130, 130]
y_z=[0,0,0,0]
vertices = [list(zip(y_x,y_y,y_z))]
poly = Poly3DCollection(vertices, alpha=0.1, color="black")
ax.add collection3d(poly)
# generate z axis
    #positive
ax.quiver(0.0, 0.0, 0.0, 0.0, 5.0, arrow length ratio=0.1, alpha = 0.5, color="black")
    #negative
ax.quiver(0.0, 0.0, 0.0, 0.0, -5.0, arrow length ratio=0.1, alpha = 0.5, color="black")
# generate z plane
\#z \ x=[0,0,0,0]
\#z \ y = [-130, 130, 130, -130]
\#z \ z = [-5.0, -5, 5.0, 5.0]
\#vertices = [list(zip(z x, z y, z z))]
#poly = Poly3DCollection(vertices, alpha=0.1, color="black")
#ax.add collection3d(poly)
ax.scatter(x_no,y_no,z_no,'o', color='red')
ax.scatter(x_yes,y_yes,z_yes,'o', color='green')
ax.scatter(x_i_x, x_i_y, x_i_z,'v', color='black')
ax.set xlabel("x:Height (cm)")
```

```
ax.set ylabel("y:Weight (kg)")
              ax.set zlabel("z:Constant 1")
              ax.quiver(0.0, 0.0, 0.0, theta[0], theta[1], theta[2], arrow length ratio=0.001, color="blue")
              perp = [190, 90, 0]
              perp[2] = -(perp[0]*theta[0]+perp[1]*theta[1])/theta[2]
              ax.quiver(0.0, 0.0, 0.0, theta[0], theta[1], theta[2], arrow length ratio=0.001, color="blue")
              ax.quiver(0.0, 0.0, 0.0, perp[0], perp[1], perp[2], arrow length ratio=0.0001, color="red")
              ax.view init(15,45)
              plt.show()
In [18]: # function that receives two vectors that can
         # be multiplied and returns their dot product
         def dotProduct(theta vector, data point x, data point y, data point z):
              result = 0
              result += theta vector[0]*data point x
              result += theta vector[1]*data point y
              result += theta vector[2]*data point z
              return result
In [19]: # formally initialize theta vector and theta 0 vector
         theta = [40, -15, -6000]
         m = 1
In [20]: # -1 Labeled red point observations
         x no = np.array([158, 165, 170, 162, 176])
         y_{no} = np.array([60, 72, 92, 65, 97])
         z no = np.array([1, 1, 1, 1, 1])
         1 \text{ no} = \text{np.array}([-1, -1, -1, -1, -1])
         # +1 labeled red point observations
         x_yes = [185, 194, 190, 197, 202]
         y \text{ yes} = [82, 87, 92, 85, 95]
         z \text{ yes} = [1, 1, 1, 1, 1]
         l_yes = np.array([1, 1, 1, 1, 1])
         # merge x-coordinates of all observations into 1 set
         x = np.append(x no, x yes)
         # merge y-coordinates of all observations into 1 set
         y = np.append(y no, y yes)
         # merge z-coordinates of all observations into 1 set
```

```
z = np.append(z no, z yes)
# merge z-coordinates of all observations into 1 set
labels = np.append(1 no, 1 yes)
# iterate from 1 to m
for t in range(1, m+1):
   # have we observed a change in theta vector
   is theta changed = False
   #iterate through data points
   for i in range(len(x)):
       print("\t0bservation data point:["+str(x[i])+","+str(y[i])+","+str(z[i])+"]")
       print("\tLabel"+str(labels[i]))
       print("\tPrediction:"+str(prediction))
       print("\tFrom Theta:"+str(theta))
       plotPerceptronResults(theta, x[i], y[i], z[i])
       prediction = labels[i] * dotProduct(theta, x[i], y[i], z[i])
       if(prediction<=0):</pre>
           print("\tWrong Prediction")
           print("\tFrom Theta:"+str(theta))
           #our hypothesis prediction is wrong and has to change
           theta[0] += labels[i]*x[i]
           theta[1] += labels[i]*y[i]
           theta[2] += labels[i]*z[i]
           is theta changed = True
           print("\tTo Theta:"+str(theta))
           print("----")
   if(is theta changed==False):
       print("Finished at:"+str(t))
       print("Theta:"+str(theta))
       plotPerceptronResults(theta, 0, 0, 0)
       break
   #print("========="")
```

Observation data point:[158,60,1]
Label-1
Prediction:41.5
From Theta:[40, -15, -6000]

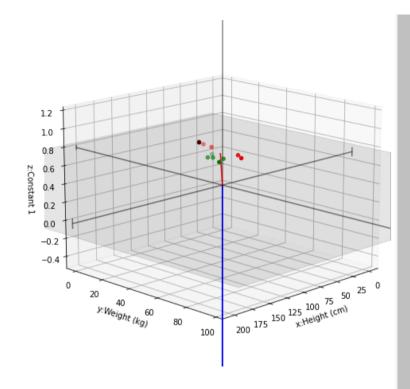


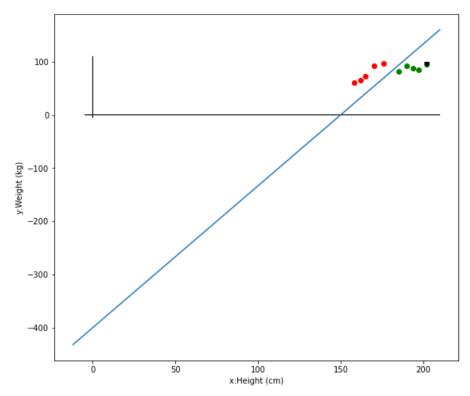
Observation data point:[165,72,1]

Label-1

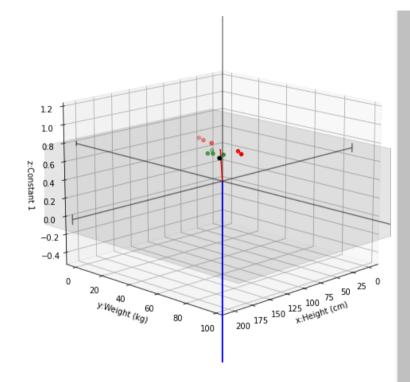
Prediction:580

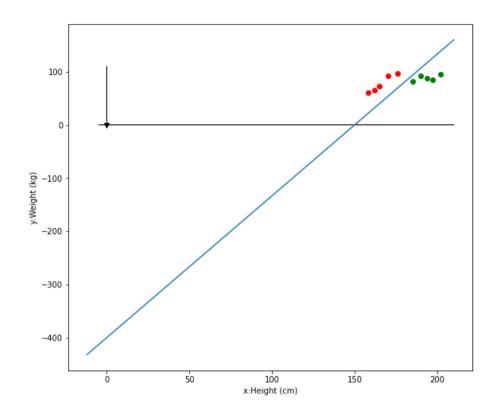
From Theta:[40, -15, -6000]

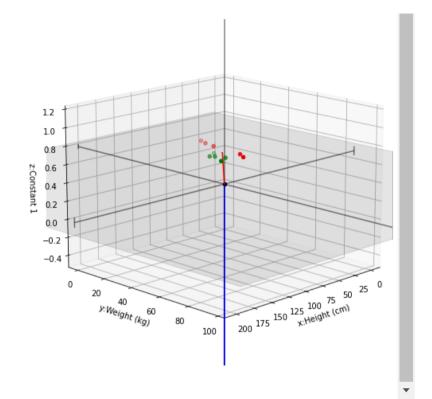




Finished at:1 Theta:[40, -15, -6000]







As we can see, for the same equation:

$$heta_1 \cdot x_1^{(i)} + heta_2 \cdot x_2^{(i)} + heta_0 \cdot x_3^{(i)} = 0$$

in the left hand size, we have the classification representation of the hypothesis that every  $x_2^{(i)}$  (or y or  $f(x_1^{(i)})$ ) must satisfy the function:

$$hypothesis: f(x_1^{(i)}) = -rac{( heta_1 \cdot x_1^{(i)} + x_3 \cdot heta_0)}{ heta_2}$$

whereas in the right hand side, we have the classification representation of the hypothesis, that must be perpendicular to  $\theta$ , meaning that for every given x, y:

$$hypothesis = \overset{-}{h} = (x,y,-rac{( heta_1\cdot x + heta_2\cdot y)}{x_3\cdot heta_0})$$

where  $\overline{h}$  passes through the origin.

Given that our perceptron algorithm is ensured to work whenever the data set is linearly classified and the classifier passes through the origin, the hypothesis vector solution in  $\mathbb{R}^3$  is exactly equivelent to the satisfaction of the hypothesis function in  $\mathbb{R}^2$ .

An additional natural interpretation that we can attribute to the hypothesis function of  $R^2$ , is that of the weighted sum, where of a given classification problem, we attribute an importance to each parameter. Namely, we can view  $\overline{\theta}$  as  $\overline{w}$  vector, where:

$$egin{aligned} heta_1 &= w_1 = importanceOf(x_1) \ heta_1 &= w_1 = importanceOf(x_2) \ heta_0 &= w_1 = bias \end{aligned}$$