#### **MAKING MODEL FOR BINARY CLASSIFICATION:**

### Classification- Breast Cancer or Not

#### **INTRODUCTION:**

- 1) ANN: Artificial neural network consists of
  - Input layer
  - Hidden layer
  - Output layer
- 2) SLP: Single layer perception
  - If ANN model has no hidden layer its called SLP
- 3) Basic equations:

#### Output = Weight \* Input + Bias

We already have output and input layers. We don't have weight (a value that can give different weights depending on features and output => [len(features), len(output)])

And bias (a value that can give different weights depending on features => [len(output)])

#### **IMPORTING LIBRARIES:**

import tensorflow as tf

import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

import seaborn as sns

from sklearn.model\_selection import train\_test\_split

from sklearn.preprocessing import MinMaxScaler, Normalizer

from sklearn.decomposition import PCA as sklearnPCA

# Supress unnecessary warnings so that presentation looks clean import warnings

warnings.filterwarnings("ignore")

#### **EXPLORING DATASET:**

1) Import data set:

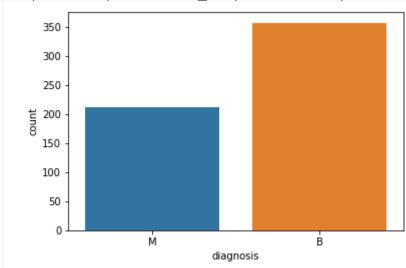
# wbcd = pd.read\_csv("../input/data.csv") wbcd.head()

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_i	
0	842302	М	17.99	10.38	122.80	1001.0	0.11840	
1	842517	М	20.57	17.77	132.90	1326.0	0.08474	
2	84300903	М	19.69	21.25	130.00	1203.0	0.10960	
3	84348301	М	11.42	20.38	77.58	386.1	0.14250	
4	84358402	М	20.29	14.34	135.10	1297.0	0.10030	
4	<b>→</b>							

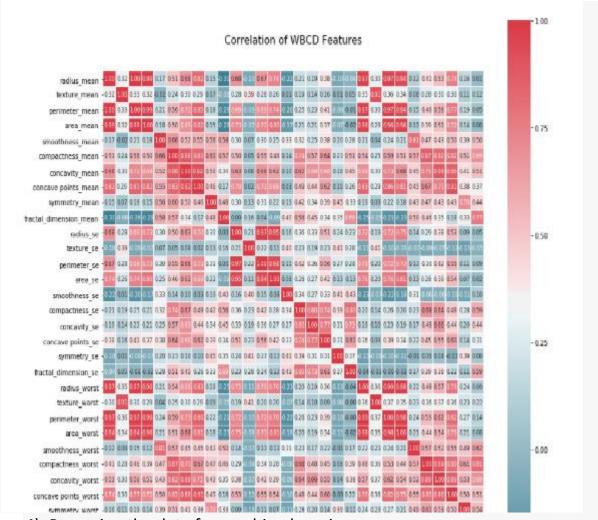
## 2) Summary the diagnosis:

INP: sns.countplot(wbcd['diagnosis'],label="Count")

Output: <matplotlib.axes.\_subplots.AxesSubplot at 0x7fece72a3470>



## 3) Correlation Plot of 30 features



4) Preparing the data for machine learning:

## i) Divide "WBCD data" into Train(70%) / Test data(30%)

```
- train,test = train_test_split(wbcd, test_size=0.3, random_state=42)
- print("Training Data :",train.shape)
- print("Testing Data :",test.shape)
```

## II) Drop ID column:

```
-train_id = train['id']

test_id = test['id']

train_data = train.iloc[:,1:]

test_data = test.iloc[:,1:]

print("Training Data :",train_data.shape)

print("Testing Data :",test_data.shape)
```

III) Seperate x:Feature data(30) / y:Result data(1)

## -Normalize x\_data values for better prediction

```
# Training Data
train_x = train_data.iloc[:,1:]
train x = MinMaxScaler().fit transform(train x)
print("Training Data :", train x.shape)
# Testing Data
test x = test data.iloc[:,1:]
test x = MinMaxScaler().fit transform(test x)
print("Testing Data :", test_x.shape)
Change Results(diagnosis) format : String -> Numeric
In [10]:
# Training Data
train_y = train_data.iloc[:,:1]
train y[train y=='M']=0
train y[train y=='B']=1
print("Training Data :", train_y.shape)
# Testing Data
test y = test data.iloc[:,:1]
test y[test y=='M']=0
test y[test y=='B']=1
print("Testing Data :", test_y.shape)
```

## **5)** Making ANN-SLP Model

1) Make "Placeholder" for dynamic variable allocation

Placeholder is one of the function in tensorflow. It is a space to put and change values while the program is running.

- for X, a place must have 30 columns, since wbcd data has 30 features.
- for Y, a place must have 1 columns, since the results has 1 outcome.

• If you see the row "None", it means it has no size limits. (You can write -1 instead of "None")

```
In [11]:
```

X = tf.placeholder(tf.float32, [None,30])

Y = tf.placeholder(tf.float32, [None, 1])

6-2) Make Weight, Bias value with randomly

- W(weight): why [30,1]? 16 for 16 features, 1 for 1 Outcome(results).
- P(weight): why [10,1]? 10 for 10 PCA features, 1 for 1 Outcome(results).
- b(bias): why [1]? outcome has 1 layers.

In [12]:

# weight

W = tf.Variable(tf.random\_normal([30,1], seed=0), name='weight')

# bias

b = tf.Variable(tf.random\_normal([1], seed=0), name='bias')
6-3) Make Output Results

- Output = Weight \* Input + Bias
- tf.matmul(): for array multiply

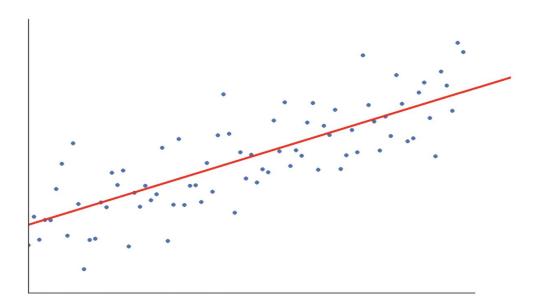
In [13]:

logits = tf.matmul(X,W) + b

6-4) Cross Entropy

Before this, you have to know How Linear Regression Works

- Linear Regression: Draw a random line to find the **mean square root error** and find the slope and intercept to minimize this value (reduce the error to the minimum)
- Since Logits is also linear equation, you have to find minimum cost!



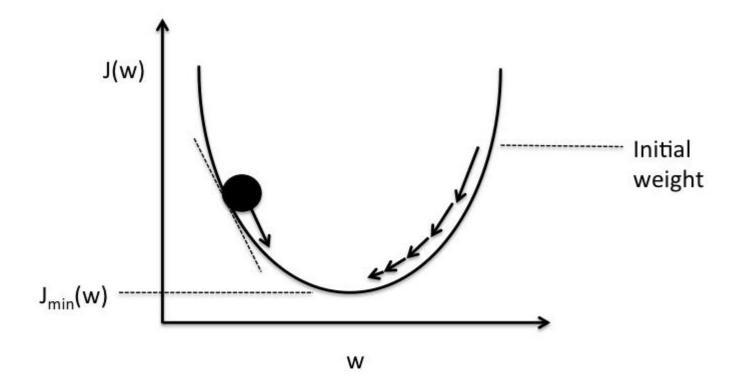
For example, logits(we get above) is **red line**, and the real dataset is **blue dot**.

- 1. For finding cost, you have to substract all blue dot value with red line.
- 2. Next, You add all distance you find and get average.
- 3. For good prediction, this average distance of red line & blue dot must be minimum value.
- 4. tf.nn.sigmoid\_cross\_entropy\_with\_logits(): for gradient\_descent with sig results(hypothesis).

```
In [14]:
```

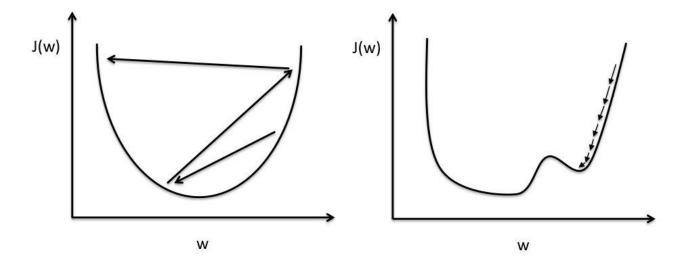
hypothesis = tf.nn.sigmoid(logits)

```
cost_i = tf.nn.sigmoid_cross_entropy_with_logits(logits=logits,labels=Y)
cost = tf.reduce_mean(cost_i)
# cost = -tf.reduce_mean(Y * tf.log(hypothesis) + (1 - Y) * tf.log(1 - hypothesis))
6-5) Gradient Descent Optimizer
```



## Schematic of gradient descent.

- GradientDescentOptimizer: It makes the best result with the least error
- There are lots of optimizer methods provided in tensorflow. (GradientDescent, Adam, RMSProp, etc.)
- learning rate: It indicates the degree of descending size.



Large learning rate: Overshooting.

Small learning rate: Many iterations until convergence and trapping in local minima.

```
In [15]:
train = tf.train.GradientDescentOptimizer(learning_rate=0.1).minimize(cost)
6-6) Compare: original vs. prediction
In [16]:
prediction = tf.cast(hypothesis > 0.5, dtype=tf.float32)
correct prediction = tf.equal(prediction, Y)
accuracy = tf.reduce_mean(tf.cast(correct_prediction, dtype=tf.float32))
6-7) Activate Model
In [17]:
with tf.Session() as sess:
  sess.run(tf.global variables initializer())
  for step in range(10001):
    sess.run(train, feed_dict={X: train_x, Y: train_y})
    if step \% 1000 == 0:
       loss, acc = sess.run([cost, accuracy], feed_dict={X: train_x, Y: train_y})
       print("Step: {:5}\tLoss: {:.3f}\tAcc: {:.2%}".format(step, loss, acc))
  train_acc = sess.run(accuracy, feed_dict={X: train_x, Y: train_y})
  test acc,test predict,test correct = sess.run([accuracy,prediction,correct pr
ediction], feed dict={X: test x, Y: test y})
  print("Model Prediction =", train acc)
  print("Test Prediction =", test acc)
```

```
5) ANN-SLP MODEL:
 train_x, test_x : normalization data
   30 features

    train y, test y

In [18]:
def ann slp():
  print("======Data Summary=======")
  print("Training Data :", train x.shape)
  print("Testing Data :", test x.shape)
  X = tf.placeholder(tf.float32, [None,30])
  Y = tf.placeholder(tf.float32, [None, 1])
  W = tf.Variable(tf.random normal([30,1], seed=0), name='weight')
  b = tf.Variable(tf.random_normal([1], seed=0), name='bias')
  logits = tf.matmul(X,W) + b
  hypothesis = tf.nn.sigmoid(logits)
  cost i = tf.nn.sigmoid cross entropy with logits(logits=logits,labels=Y)
  cost = tf.reduce_mean(cost_i)
  train = tf.train.GradientDescentOptimizer(learning rate=0.1).minimize(cost)
  prediction = tf.cast(hypothesis > 0.5, dtype=tf.float32)
  correct prediction = tf.equal(prediction, Y)
  accuracy = tf.reduce mean(tf.cast(correct prediction, dtype=tf.float32))
  print("\n===========")
  with tf.Session() as sess:
    sess.run(tf.global_variables_initializer())
    for step in range(10001):
      sess.run(train, feed dict={X: train x, Y: train y})
      if step \% 1000 == 0:
        loss, acc = sess.run([cost, accuracy], feed dict={X: train x, Y: train y})
        print("Step: {:5}\tLoss: {:.3f}\tAcc: {:.2%}".format(step, loss, acc))
    train_acc = sess.run(accuracy, feed_dict={X: train_x, Y: train_y})
```

```
test_acc,test_predict,test_correct = sess.run([accuracy,prediction,correct_prediction], feed_dict={X: test_x, Y: test_y})

print("\n========Results======")
print("Model Prediction =", train_acc)
print("Test Prediction =", test_acc)

return train_acc,test_acc

ann_slp_train_acc, ann_slp_test_acc = ann_slp()
```