

## Statistical Analysis of Breast Cancer Data and EDA

```
%%HTML
<script src="./require.js"></script>

<IPython.core.display.HTML object>

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

# Importing data
data = pd.read_csv('./Breast Cancer Detection.csv')
del data['Unnamed: 32']
```

### Performing Data Wrangling Operations

```
X = data.iloc[:, 2:].values
y = data.iloc[:, 1].values
```

Replacing Null Values if any.

```
data.isnull().sum()

id                0
diagnosis         0
radius_mean       0
texture_mean      0
perimeter_mean    0
area_mean         0
smoothness_mean   0
compactness_mean  0
concavity_mean    0
concave points_mean 0
symmetry_mean     0
fractal_dimension_mean 0
radius_se         0
texture_se        0
perimeter_se      0
area_se           0
smoothness_se     0
compactness_se    0
concavity_se      0
concave points_se 0
symmetry_se       0
fractal_dimension_se 0
radius_worst      0
```

```

texture_worst      0
perimeter_worst    0
area_worst         0
smoothness_worst   0
compactness_worst  0
concavity_worst    0
concave points_worst 0
symmetry_worst     0
fractal_dimension_worst 0
dtype: int64

```

Total number of counts for Malignant (M) and Benign (B)

```

data['diagnosis'].value_counts()

diagnosis
B      357
M      212
Name: count, dtype: int64

```

A quick description of data

```

data.describe()

          id  radius_mean  texture_mean  perimeter_mean
area_mean
count  5.690000e+02    569.000000    569.000000    569.000000
569.000000 \
mean   3.037183e+07    14.127292    19.289649    91.969033
654.889104
std    1.250206e+08    3.524049    4.301036    24.298981
351.914129
min    8.670000e+03    6.981000    9.710000    43.790000
143.500000
25%    8.692180e+05    11.700000    16.170000    75.170000
420.300000
50%    9.060240e+05    13.370000    18.840000    86.240000
551.100000
75%    8.813129e+06    15.780000    21.800000    104.100000
782.700000
max    9.113205e+08    28.110000    39.280000    188.500000
2501.000000

          smoothness_mean  compactness_mean  concavity_mean  concave
points_mean
count      569.000000    569.000000    569.000000
569.000000 \
mean        0.096360        0.104341        0.088799
0.048919

```

std	0.014064	0.052813	0.079720
0.038803			
min	0.052630	0.019380	0.000000
0.000000			
25%	0.086370	0.064920	0.029560
0.020310			
50%	0.095870	0.092630	0.061540
0.033500			
75%	0.105300	0.130400	0.130700
0.074000			
max	0.163400	0.345400	0.426800
0.201200			

	symmetry_mean	...	radius_worst	texture_worst
perimeter_worst				
count	569.000000	...	569.000000	569.000000
569.000000	\			
mean	0.181162	...	16.269190	25.677223
107.261213				
std	0.027414	...	4.833242	6.146258
33.602542				
min	0.106000	...	7.930000	12.020000
50.410000				
25%	0.161900	...	13.010000	21.080000
84.110000				
50%	0.179200	...	14.970000	25.410000
97.660000				
75%	0.195700	...	18.790000	29.720000
125.400000				
max	0.304000	...	36.040000	49.540000
251.200000				

	area_worst	smoothness_worst	compactness_worst
concavity_worst			
count	569.000000	569.000000	569.000000
569.000000	\		
mean	880.583128	0.132369	0.254265
0.272188			
std	569.356993	0.022832	0.157336
0.208624			
min	185.200000	0.071170	0.027290
0.000000			
25%	515.300000	0.116600	0.147200
0.114500			
50%	686.500000	0.131300	0.211900
0.226700			
75%	1084.000000	0.146000	0.339100
0.382900			
max	4254.000000	0.222600	1.058000

1.252000

	concave	points_worst	symmetry_worst	fractal_dimension_worst
count	569.000000	569.000000	569.000000	569.000000
mean	0.114606	0.290076	0.083946	0.083946
std	0.065732	0.061867	0.018061	0.018061
min	0.000000	0.156500	0.055040	0.055040
25%	0.064930	0.250400	0.071460	0.071460
50%	0.099930	0.282200	0.080040	0.080040
75%	0.161400	0.317900	0.092080	0.092080
max	0.291000	0.663800	0.207500	0.207500

[8 rows x 31 columns]

Categorical Observation to Detect Breast Cancer across Radius, Perimeter, Area, Smoothness, Concavity and Worst Fractal Dimensions.

```
import plotly.express as px
import plotly.io as pio
pio.renderers.default='notebook'

px.scatter_matrix(data, dimensions=["radius_mean", "perimeter_mean",
"area_mean", "smoothness_mean", "concavity_mean",
"fractal_dimension_worst"], color="diagnosis", title="Malignancy
Analysis for Given Features.")
```

Exploring Distribution of Cases over Fractal Dimensions. (M = Malignant, B = Benign)

```
fig = px.box(data, x='diagnosis', y= 'fractal_dimension_worst')
fig.show()
```

Using the data to create a Deep Learning Model to predict breast cancer with given features.

```
# Encoding categorical data
from sklearn.preprocessing import LabelEncoder
labelencoder_X_1 = LabelEncoder()
y = labelencoder_X_1.fit_transform(y)

# Splitting the dataset into the Training set and Test set
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size =
0.1, random_state = 0)

#Feature Scaling
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)
```

Using Keras and Tensorflow package to develop CNN model

```
import keras
from keras.models import Sequential
from keras.layers import Dense, Dropout

# Initialising the ANN
classifier = Sequential()
```

Developing First Input Layer using RELU activation

```
# Adding the input layer and the first hidden layer
classifier.add(Dense(16, activation='relu'))
# Adding dropout to prevent overfitting
classifier.add(Dropout(0.1))
```

Developing Second Input Layer Using RELU Activation

```
# Adding the second hidden layer
classifier.add(Dense(16, activation='relu'))
# Adding dropout to prevent overfitting
classifier.add(Dropout(0.1))
```

Using sigmoid function as gradient descent algorithm for backward propagation.

```
classifier.add(Dense(1, activation='sigmoid'))
```

Optimizing the Classifier with Adam algorithm and reducing loss rate with Binary Crossentropy.

```
classifier.compile(optimizer='adam', loss='binary_crossentropy',
metrics=['accuracy'])
```

Fitting the classifier with training sets.

```
classifier.fit(X_train, y_train, batch_size=100)

6/6 [=====] - 0s 2ms/step - loss: 0.1987 -
accuracy: 0.9551

<keras.callbacks.History at 0x14804c42740>
```

Running iterations to optimize prediction accuracy

```
# Predicting the Test set results
y_pred = classifier.predict(X_test)
y_pred = (y_pred > 0.5)

2/2 [=====] - 0s 2ms/step
```

```
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test, y_pred)

print("Our accuracy is {}".format(((cm[0][0] + cm[1][1])/57)*100))

Our accuracy is 91.22807017543859%

sns.heatmap(cm,annot=True)

<Axes: >
```

