

About Problem Statement and **Data Set Exploratory** Data Analysis

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About Data Set

- 1. ID number
- 2. Diagnosis (M = malignant, B = benign)
- 3. Ten real-valued features are computed for each cell nucleus:

Problem Statement

- Its a Breast Cancer Classification Problem.
- Breast cancer is the most common cancer amongst women in the world. It accounts for 25% of all cancer cases
- These cells usually form tumors that can be seen via X-ray or felt as lumps in the breast area.
- The key challenges against it's detection is how to classify tumors into malignant (cancerous) or benign(non cancerous).

Data Source : https://www.k

- radius (mean of distances from center to points on the perimeter)
- 2. texture (standard deviation of gray-scale values)
- 3. perimeter
- 4. area
- 5. smoothness (local variation in radius lengths)
- 6. compactness (perimeter^2 / area 1.0)
- 7. concavity (severity of concave portions of the contour)
- 8. concave points (number of concave portions of the contour)
- 9. symmetry
- 10. fractal dimension ("coastline approximation" 1)

The mean, standard error and "worst" or largest (mean of the three largest values) of these features were computed for resulting in 30 features. For instance, field 3 is Mean Radius, field 13 is Radius SE, field 23 is Worst Radius.

EXPLORATORY DATA ANALYSIS radius mean texture mean 10 35 40 | 10 I perimeter mean area mean 20 20 100 125 150 175 1000 1500 2500 2000 smoothness mean compactness mean 30 В 20 20 10 10 0.06 0.08 0.10 0.12 0.14 0.16 0.05 0.10 0.15 0.20 0.25 0.30 0.35 concavity_mean concave points mean В 20 20 0.2 0.3 0.05 0.10 0.15 0.0 0.4 0.00 0.20 symmetry_mean fractal dimension mean 30 30 20 -20 10 10 0.05 0.10 0.15 0.20 0.25 0.30 0.06 0.07 0.08 0.09

MEAN OF FEATURES



It can be observed that Larger the mean values of radius", concave points", concavity", "compactness, area and perimeter" shows correlation with Malignant tumors.

So these features can be used for classification.

On the other hand mean values of texture, smoothness, symmetry or fractual dimension does not show a particular preference of one diagnosis over the other

EXPLORATORY DATA ANALYSIS radius_se texture se в 50 > 25 1.0 2.5 0.0 perimeter se area se 100 150 100 50 50 20 100 200 300 400 500 smoothness_se compactness_se 40 20 20 0.005 0.010 0.015 0.020 0.025 0.030 0.00 0.02 0.04 0.06 0.08 0.10 0.12 0.14 concave points_se concavity_se 75 50 20 25 0.4 0.0 0.2 0.3 0.00 0.02 0.03 0.04 0.05 0.1 0.01 symmetry_se fractal_dimension_se 40 40 20 20 0.01 0.02 0.03 0.04 0.05 0.06 0.07 0.08 0.000 0.005 0.010 0.015 0.020 0.025 0.030

STANDARD ERROR OF ATTRIBUTES



* It can be observed that Larger the standard error values of radius", concave points", "compactness, area and perimeter" shows correlation with Malignant tumors. So these features can be used for classification.

EXPLORATORY DATA ANALYSIS radius_worst texture worst 20 30 perimeter_worst area worst 20 25 250 200 2000 3000 4000 smoothness worst compactness_worst 30 20 20 10 10 0.075 0.100 0.125 0.150 0.175 0.200 0.225 0.6 concave points_worst concavity_worst 20 0.6 0.8 1.0 0.00 0.05 0.10 0.15 0.20 0.25 0.30 1.2 symmetry_worst fractal dimension worst 30 20 20 10 0.050 0.075 0.100 0.125 0.150 0.175 0.200 0.6

WORST/LARGEST VALUE OF ATTRIBUTES



- * Similar to means values, It can be observed that Larger the worst values of radius", concave points", concavity", "compactness, area and perimeter" shows correlation with Malignant tumors. So these features can be used for classification.
- * On the other hand worst values of texture, smoothness, symmetry or fractual dimension does not show a particular preference of one diagnosis over the other

```
data_imp = data[important_feats]
target = data["diagnosis"]
```

Data Preprocessing

- New Dataset of Important features is created.
- Dependent and independent variables are seperated.
- Independent variables are **Normalized** using Min Max Scalar .
- Splitting the data into train and test data.

```
# normalising data
x = MinMaxScaler().fit_transform(data_imp)
y = np.array(target)

# splitting data
x_train , x_test , y_train , y_test = train_test_split(x,y, train_size = 0.2)
```

Random Forest Classification Report

Predictions and ML Models

- I have trained data on 2 different machine learning models Random Forest classifier and SVM
- Out of 2 Random Forest Classifier outperformed with 94% Accuracy , 95 % precision and 96% recall.

		precision	recall	f1-score	support
	0	0.95	0.96	0.95	290
	1	0.92	0.92	0.92	166
accurac	y			0.94	456
macro av	g	0.94	0.94	0.94	456
weighted av	g	0.94	0.94	0.94	456

	precision	recall	f1-score	support
0	0.92	0.98	0.95	290
1	0.97	0.85	0.90	166
accuracy			0.93	456
macro avg	0.94	0.92	0.93	456
weighted avg	0.94	0.93	0.93	456
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SVM Classification Report