

Python Final Group Project

“Breast Cancer Detection”

Class: Python Programming (CS-661)

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Group 8 Members:

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1. Topic

: “Breast Cancer Detection”

2. Summary

- Dataset: Kaggle

Dataset : <https://www.kaggle.com/code/faresmohammad/breast-cancer-detection>

- Purpose:

The purpose of our project is to predict with accuracy if a tumor cell is malignant or benign by utilizing machine learning algorithms with various predictive models for breast cancer detection. By employing different classification algorithms, we aim to explore and compare their performance in accurately identifying whether a tumor is malignant or benign based on the provided attributes.

-Independent Variable (X) = All the columns except for the 'Diagnosis' variable

-Dependent Variable(y) = 'Diagnosis' (Predicting if a patient has cancer (1=M) or not (0=B))

The analysis involved data preprocessing, exploratory analysis, outlier detection, model building, evaluation, and comparison. Each step contributes to understanding the dataset and evaluating the performance of various machine learning algorithms in breast cancer detection.

1. Exploratory Analysis: Conduct exploratory data analysis (EDA) to understand the distribution and relationships between different features within the dataset. This will involve visualizations and statistical summaries to gain insights into the data.

2. Feature Importance: Evaluate the importance of features in determining the nature of the tumor (malignant or benign) using various algorithms such as Decision Trees, Random Forests, and others to identify which attributes have the most significant impact on the classification.

3. Model Development: Implement different classification algorithms including Logistic Regression, Naive Bayes, Support Vector Machines (SVM), K-Nearest Neighbors (KNN), Decision Trees, and Random Forests. Compare and evaluate their performance in terms of accuracy, precision, recall, F1-score, and area under the ROC curve.

4. Model Comparison: Analyze and compare the performance metrics of each model to identify the most effective algorithm for breast cancer detection based on the dataset.

5. Optimization and Validation: Optimize the selected models by tuning hyperparameters and conduct cross-validation to ensure robustness and reliability of the chosen algorithm(s) for accurate prediction.

3. Predictive Models

1. EDA - Lolyna / 2. Logistic Regression (Scikit-learn) - Ebukka/

3. Support Vector Machine (Linear) (Scikit-learn) - Abhay / 4. K-Nearest Neighbours (Scikit-learn) - Sunny

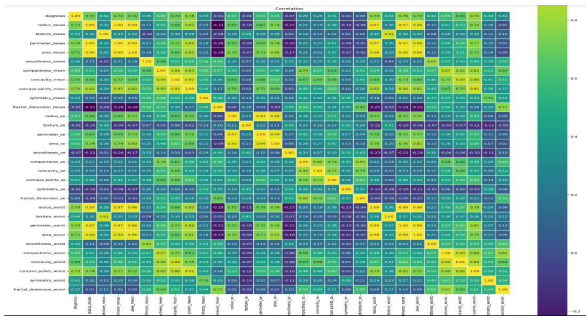
5. Decision Tree (Scikit-learn) - Ibrahim/ 6. Random Forest (Scikit-learn) - Niyati

4. Data Visualization

```
#Describe the database  
df.describe()
```

| | diagnosis | radius_mean | texture_mean | perimeter_mean |
|-------|------------|-------------|--------------|----------------|
| count | 569.000000 | 569.000000 | 569.000000 | 569.000000 |
| mean | 0.372583 | 14.127292 | 19.289649 | 91.969033 |
| std | 0.483918 | 3.524049 | 4.301036 | 24.298981 |
| min | 0.000000 | 6.981000 | 9.710000 | 43.790000 |
| 25% | 0.000000 | 11.700000 | 16.170000 | 75.170000 |
| 50% | 0.000000 | 13.370000 | 18.840000 | 86.240000 |
| 75% | 1.000000 | 15.780000 | 21.800000 | 104.100000 |
| max | 1.000000 | 28.110000 | 39.280000 | 188.500000 |

8 rows x 31 columns



Accuracy of model
62.28070175438597 % Logistic Regression

Accuracy: 0.9385964912280702 Decision Tree Model
Classification Report:

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| B | 0.96 | 0.94 | 0.95 | 71 |
| M | 0.91 | 0.93 | 0.92 | 43 |
| accuracy | | | 0.94 | 114 |
| macro avg | 0.93 | 0.94 | 0.93 | 114 |
| weighted avg | 0.94 | 0.94 | 0.94 | 114 |

```
# classification report  
target_names = ['Patient has Cancer', 'Patient DOES NOT have Cancer']  
print('Classification report: \n', classification_report(y_test, y_pred))
```

Classification report:

| | precision | recall | f1-score | support |
|------------------------------|-----------|--------|----------|---------|
| Patient has Cancer | 0.62 | 1.00 | 0.77 | 71 |
| Patient DOES NOT have Cancer | 0.00 | 0.00 | 0.00 | 43 |
| accuracy | | | 0.62 | 114 |
| macro avg | 0.31 | 0.50 | 0.38 | 114 |
| weighted avg | 0.39 | 0.62 | 0.48 | 114 |

The accuracy score: 96.49 Random Forest
The predict score achieved using Random Forest is:

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| B | 0.96 | 0.99 | 0.97 | 71 |
| M | 0.98 | 0.93 | 0.95 | 43 |
| accuracy | | | 0.96 | 114 |
| macro avg | 0.97 | 0.96 | 0.96 | 114 |
| weighted avg | 0.97 | 0.96 | 0.96 | 114 |

Support Vector Machine Algorithm

K-NN/ Accuracy: 81.82%

Accuracy: 0.6667
Precision: 0.6667
Recall: 2.0000
Specificity: 0.6667

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| B | 0.80 | 0.97 | 0.87 | 93 |
| M | 0.90 | 0.54 | 0.68 | 50 |
| accuracy | | | 0.82 | 143 |
| macro avg | 0.85 | 0.75 | 0.77 | 143 |
| weighted avg | 0.83 | 0.82 | 0.80 | 143 |