Diagnosing Breast Cancer using Julia

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Problem Description

The problem is to diagnose the breast cancer.

- Early diagnosing breast cancer can promote timely clinical treatment to patients.
- Correct classification of the benign tumor can prevent patients from undergoing unnecessary treatments.

Thus, diagnosing breast cancer and accurate prediction of malignant and benign tumors is an important.

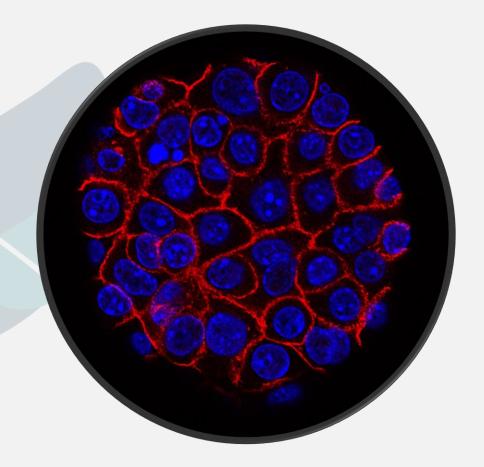
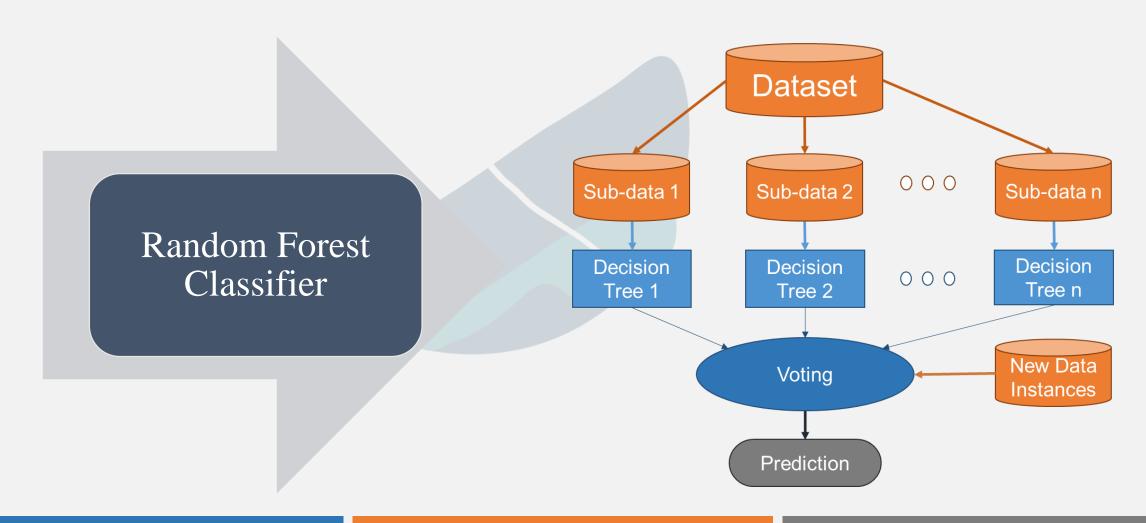


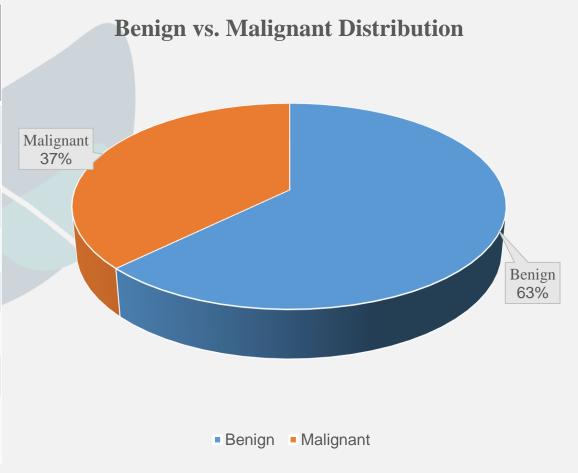
Image Source: USC Norris Comprehensive Cancer Center

Classification Method



Dataset Description

Dataset Name	Breast Cancer Wisconsin(Diagnostic) Dataset	
Source	<u>Kaggle</u>	
Number of Instances	569	
Features	31	
Feature Type	Numeric and Categorical	
Missing Values	No	
Class Value	2	
Learning Type	Classification	
Class Instances	Benign => 357 Malignant => 212	



Performance Evaluation Criteria

$$ightharpoonup$$
 Accuracy = $\frac{TP+TN}{TP+FP+FN+TN}$

> **TP rate or Recall** =
$$\frac{TP}{TP+FN}$$

$$ightharpoonup \mathbf{FP}$$
 rate = $\frac{FP}{FP+TN}$

> TN rate or Specificity =
$$\frac{TN}{FP+TN}$$

$$ightharpoonup$$
 FNrate = $\frac{FN}{FN+TP}$

$$ightharpoonup$$
 Precision = $\frac{TP}{TP+FN}$

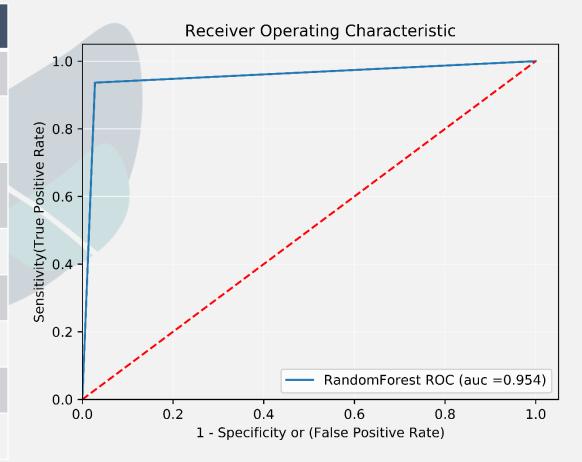
> **F1_score** =
$$\frac{2 \times TP}{2 \times TP + FP + FN}$$

$$ightharpoonup AUC = \frac{TP \text{rate} + TN \text{rate}}{2}$$

$$ightharpoonup Gmean = \sqrt{TPrate + TNrate}$$

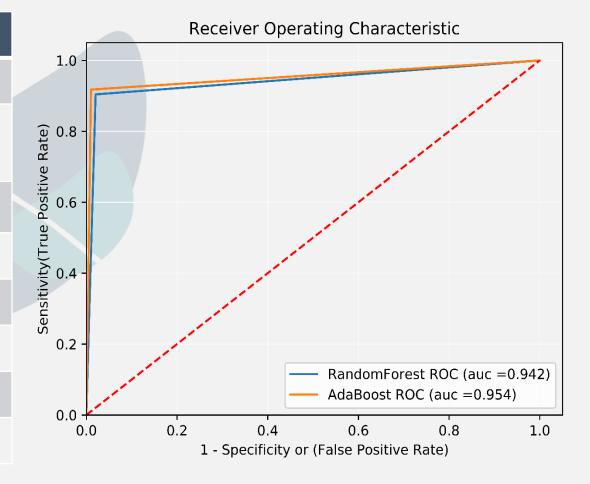
Results

Performance Metric	Score
Accuracy	96.10%
True Positive Rate or Sensitivity or Recall	92.96%
Specificity or True Negative Rate	98.09%
AUC	95.54%
Gmean	95.49%
Precision	96.81%
F1_score	94.83%
False Positive Rate	01.90%



Results

Metrics	Random Forest	AdaBoost
Accuracy	0.961014	0.964912
True Positive Rate or Sensitivity or Recall	0.929631	0.933741
Specificity or True Negative Rate	0.980988	0.98439
AUC	0.95531	0.959065
Gmean	0.954923	0.958665
Precision	0.968115	0.973164
F1_score	0.948361	0.952829
False Positive Rate	0.019012	0.01561



Julia Packages

- DataFrames
- > CSV
- Random
- FreqTables
- > PyPlot
- MLLabelUtils
- MLDataUtils
- DecisionTree

- > ScikitLearn
- > PyCall
- > MLBase

Any Question?

Thank You