

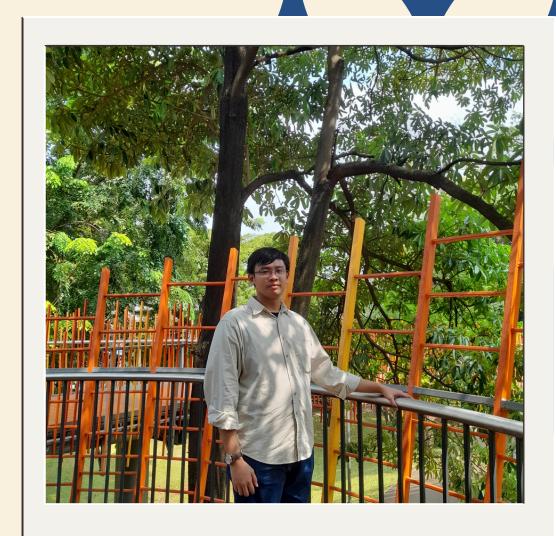
### Simple Classification of Wisconsin Breast Cancer Diagnostic Using Random Forest Algorithm

Danica Alana Sjurjahady

### ABOUT ME

Fresh graduate majoring in Agro-Industrial Technology from University of Darussalam Gontor, who have a strong interest in data analysis. Have a good understanding of basic concepts of statistics and machine learning.

Skilled in using MySQL, Python, Google Looker Studio, Google Colab, and Microsoft Power Bl. Ready to learn and grow in the role of Data Analyst.



# INTRODUCTION & OBJECTIVE

Wisconsin Breast Cancer Diagnostic is one of the toy datasets from scikit-learn. It's a classic and very easy binary classification dataset. It has 569 instances, and attributes; including: 30 numeric, predictive attributes and the class.

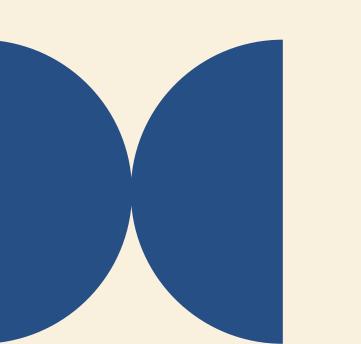
The objective is to build a machine learning model for predicting whether a breast tumor is benign (0) or malignant (1) using Random Forest Algorithm.

#### TOOLS USED

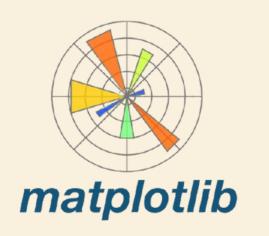












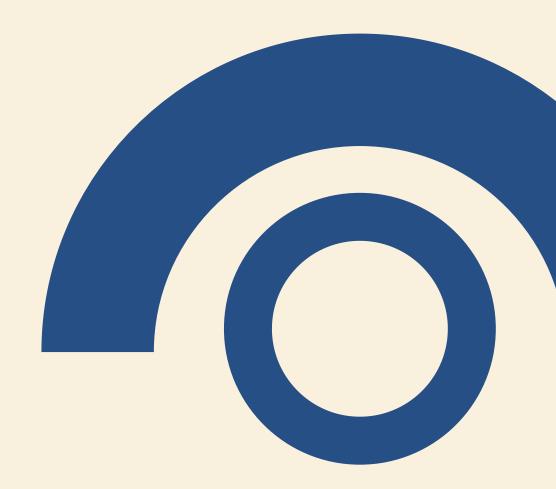
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Data
Modelling
Visualization





# INPUT DATA

```
[1] import pandas as pd
    from sklearn import datasets

# Load the Wine dataset from scikit-learn and convert it to a DataFrame
    breast_cancer = datasets.load_breast_cancer()

x = breast_cancer.data  # inputs for machine learning
y = breast_cancer.target # desired output of machine learning

# Convert feature and target data into a DataFrame
    df_x = pd.DataFrame(x, columns = breast_cancer.feature_names)
    df_y = pd.Series(y, name = 'target')

# Combine features and targets in one DataFrames
    df = pd.concat([df_x, df_y], axis = 1)

    df.head(10)
```

<b></b> _	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	 worst texture	worst perimeter	worst area	worst smoothness	worst compactness	worst concavity	worst concave points	worst symmetry	worst fractal dimension	target
C	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.30010	0.14710	0.2419	0.07871	17.33	184.60	2019.0	0.1622	0.6656	0.7119	0.2654	0.4601	0.11890	0
1	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.08690	0.07017	0.1812	0.05667	23.41	158.80	1956.0	0.1238	0.1866	0.2416	0.1860	0.2750	0.08902	0
2	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.19740	0.12790	0.2069	0.05999	25.53	152.50	1709.0	0.1444	0.4245	0.4504	0.2430	0.3613	0.08758	0
3	11.42	20.38	77.58	386.1	0.14250	0.28390	0.24140	0.10520	0.2597	0.09744	26.50	98.87	567.7	0.2098	0.8663	0.6869	0.2575	0.6638	0.17300	0
4	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.19800	0.10430	0.1809	0.05883	16.67	152.20	1575.0	0.1374	0.2050	0.4000	0.1625	0.2364	0.07678	0
5	12.45	15.70	82.57	477.1	0.12780	0.17000	0.15780	0.08089	0.2087	0.07613	23.75	103.40	741.6	0.1791	0.5249	0.5355	0.1741	0.3985	0.12440	0
6	18.25	19.98	119.60	1040.0	0.09463	0.10900	0.11270	0.07400	0.1794	0.05742	27.66	153.20	1606.0	0.1442	0.2576	0.3784	0.1932	0.3063	0.08368	0
7	13.71	20.83	90.20	577.9	0.11890	0.16450	0.09366	0.05985	0.2196	0.07451	28.14	110.60	897.0	0.1654	0.3682	0.2678	0.1556	0.3196	0.11510	0
8	13.00	21.82	87.50	519.8	0.12730	0.19320	0.18590	0.09353	0.2350	0.07389	30.73	106.20	739.3	0.1703	0.5401	0.5390	0.2060	0.4378	0.10720	0
g	12.46	24.04	83.97	475.9	0.11860	0.23960	0.22730	0.08543	0.2030	0.08243	40.68	97.65	711.4	0.1853	1.0580	1.1050	0.2210	0.4366	0.20750	0

# EXPLORATORY DATA ANALYSIS (EDA)

# View basic information about the data
 df.info()

<<class 'pandas.core.frame.DataFrame'>
 RangeIndex: 569 entries, 0 to 568
 Data columns (total 31 columns):

memory usage: 137.9 KB

Data	columns (total 31 columns	5):	
#	Column	Non-Null Count	Dtype
0	mean radius	569 non-null	float64
1	mean texture	569 non-null	float64
2	mean perimeter	569 non-null	float64
3	mean area	569 non-null	float64
4	mean smoothness	569 non-null	float64
5	mean compactness	569 non-null	float64
6	mean concavity	569 non-null	float64
7	mean concave points	569 non-null	float64
8	mean symmetry	569 non-null	float64
9	mean fractal dimension	569 non-null	float64
10	radius error	569 non-null	float64
11	texture error	569 non-null	float64
12	perimeter error	569 non-null	float64
13	area error	569 non-null	float64
14	smoothness error	569 non-null	float64
15	compactness error	569 non-null	float64
16	concavity error	569 non-null	float64
17	concave points error	569 non-null	float64
18	symmetry error	569 non-null	float64
19	fractal dimension error	569 non-null	float64
20	worst radius	569 non-null	float64
21	worst texture	569 non-null	float64
22	worst perimeter	569 non-null	float64
23	worst area	569 non-null	float64
24	worst smoothness	569 non-null	float64
25	worst compactness	569 non-null	float64
26	worst concavity	569 non-null	float64
27	worst concave points	569 non-null	float64
28	worst symmetry	569 non-null	float64
29	worst fractal dimension	569 non-null	float64
30	target	569 non-null	int64
dtype	es: float64(30), int64(1)		

[3] # Identify all the different numbers that appear in the 'target' column
 df['target'].unique()

→ array([0, 1])

[4] # View a statistical description of the data
 df.describe()

2		mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	 worst texture	ا peri
	count	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.00
	mean	14.127292	19.289649	91.969033	654.889104	0.096360	0.104341	0.088799	0.048919	0.181162	0.062798	25.677223	107.20
	std	3.524049	4.301036	24.298981	351.914129	0.014064	0.052813	0.079720	0.038803	0.027414	0.007060	6.146258	33.60
	min	6.981000	9.710000	43.790000	143.500000	0.052630	0.019380	0.000000	0.000000	0.106000	0.049960	12.020000	50.4 <sup>-</sup>
	25%	11.700000	16.170000	75.170000	420.300000	0.086370	0.064920	0.029560	0.020310	0.161900	0.057700	21.080000	84.1 <sup>-</sup>
	50%	13.370000	18.840000	86.240000	551.100000	0.095870	0.092630	0.061540	0.033500	0.179200	0.061540	25.410000	97.60
	75%	15.780000	21.800000	104.100000	782.700000	0.105300	0.130400	0.130700	0.074000	0.195700	0.066120	29.720000	125.40
	max	28.110000	39.280000	188.500000	2501.000000	0.163400	0.345400	0.426800	0.201200	0.304000	0.097440	49.540000	251.20
	8 rows ×	31 columns											

## DATA MODELLING

```
[5] from sklearn.model_selection import train_test_split

# Split the data into train and test
x_train, x_test, y_train, y_test = train_test_split(df_x, df_y, test_size = 0.2, random_state = 42)
```

[6] from sklearn.ensemble import RandomForestClassifier

# Create and train a Random Forest model
model = RandomForestClassifier(n\_estimators=100, random\_state=42)
model.fit(x\_train, y\_train)

\*\* RandomForestClassifier

\*\* RandomForestClassifier(random\_state=42)

```
[7] from sklearn.metrics import accuracy_score, classification_report

# Predict and evaluate the model
y_pred = model.predict(x_test)

accuracy = accuracy_score(y_test, y_pred)

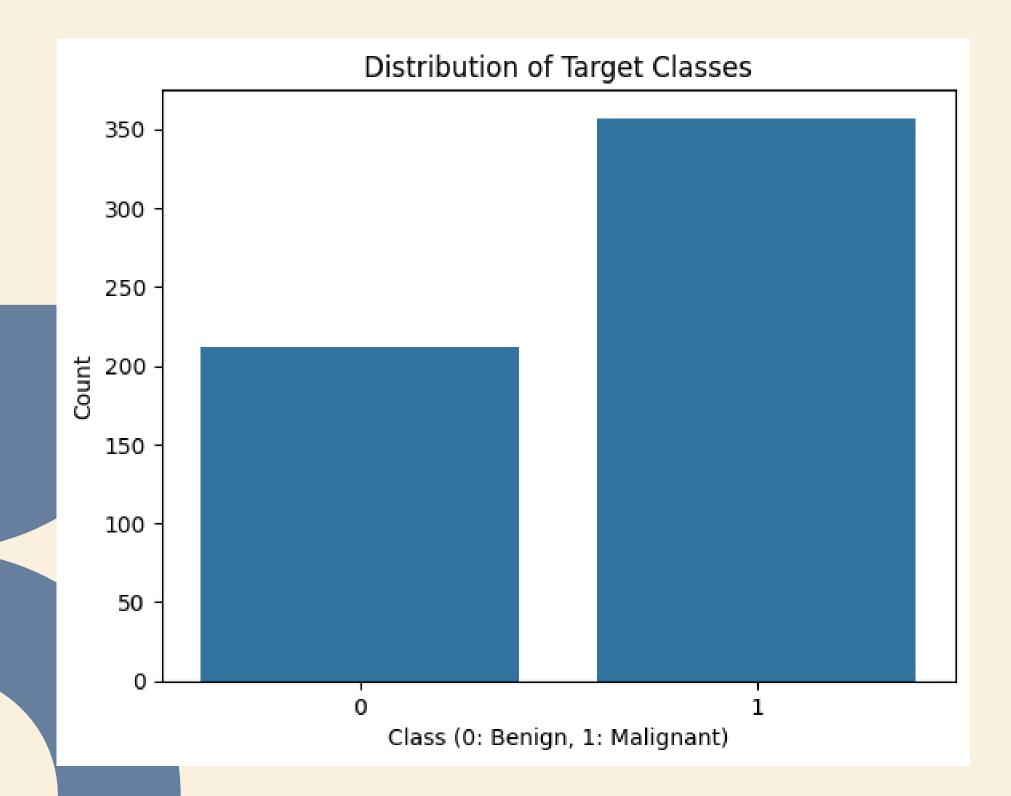
print("Classification Report:")
print(classification_report(y_test, y_pred))
print(f"Accuracy: {accuracy * 100:.2f}%")
```

```
→ Classification Report:
                              recall f1-score support
                 precision
                                0.93
                                          0.95
                      0.98
                                                     43
                      0.96
                                0.99
                                          0.97
                                                     71
                                          0.96
                                                    114
        accuracy
                                          0.96
       macro avg
                      0.97
                                0.96
                                                    114
    weighted avg
                      0.97
                                0.96
                                          0.96
                                                    114
    Accuracy: 96.49%
```

Distribution of Target Classes

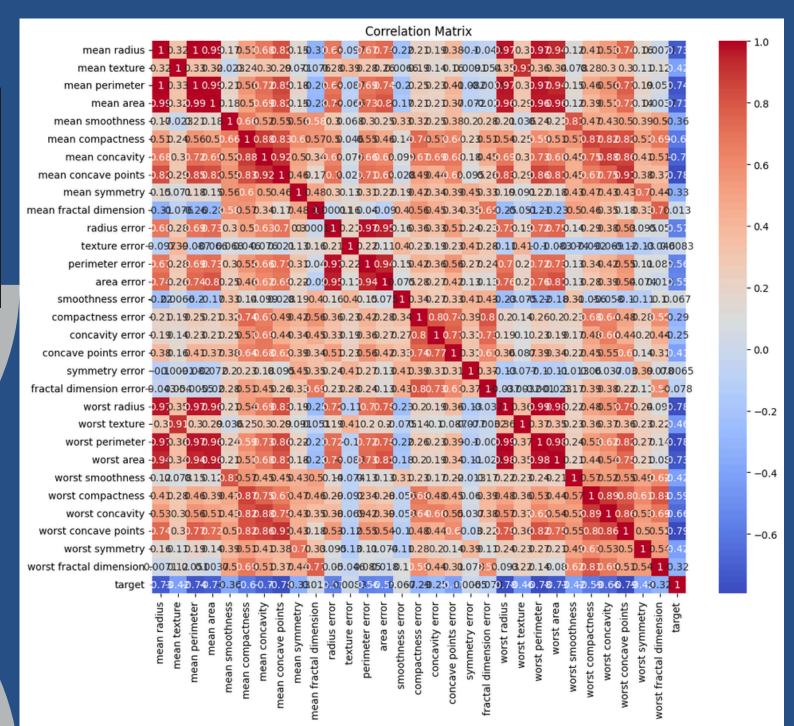
```
[8] import matplotlib.pyplot as plt
import seaborn as sns

# Visualize the distribution of target classes
sns.countplot(x='target', data=df)
plt.title('Distribution of Target Classes')
plt.xlabel('Class (0: Benign, 1: Malignant)')
plt.ylabel('Count')
plt.show()
```



#### **Correlation Matrix**

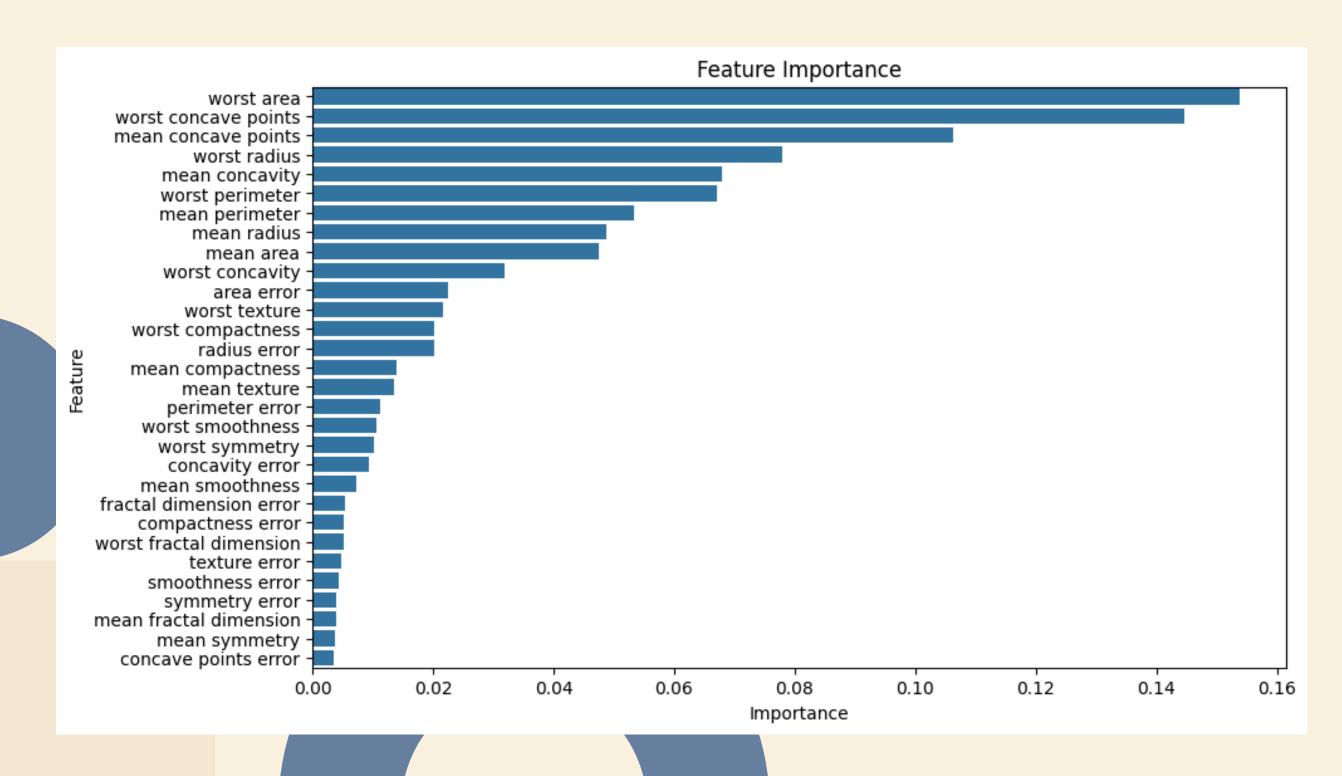
```
[9] # Visualize the correlation matrix
    plt.figure(figsize=(12, 10))
    sns.heatmap(df.corr(), annot=True, cmap='coolwarm')
    plt.title('Correlation Matrix')
    plt.show()
```



#### Feature Importance

```
[10] # Visualize feature importance from the Random Forest model
     importances = model.feature_importances_
     # Access feature names from the breast_cancer dataset's feature_names attribute
     feature_names = breast_cancer.feature_names
     feature_importance_df = pd.DataFrame({'Feature': feature_names, 'Importance': importances})
     feature_importance_df = feature_importance_df.sort_values(by='Importance', ascending=False)
     plt.figure(figsize=(10, 6))
     sns.barplot(x='Importance', y='Feature', data=feature_importance_df)
     plt.title('Feature Importance')
     plt.xlabel('Importance')
     plt.ylabel('Feature')
     plt.show()
```

#### Feature Importance



# THANK YOU

If you have any questions, suggestions or feedbacks, please do not hesitate to reach me through the contacts below:



rsgame99@gmail.com



github.com/DanicaAlana



linkedin.com/in/danica-alanasjurjahady-85b124211/

