

# LAB NO 03

## 1. Import Required Libraries

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

from sklearn.datasets import load_breast_cancer
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
```

## 2. Load Dataset

- Use a dataset Breast Cancer Dataset from sklearn.datasets.

```
data = load_breast_cancer()
df = pd.DataFrame(data.data, columns=data.feature_names)
df['target'] = data.target
```

## 3. Exploratory Data Analysis (EDA)

- Display first few rows of the dataset.
- Check for missing values.
- Plot histograms or distribution of features.
- Check class distribution (malignant vs. benign).

```
[ ] print(df.head())
```

```
▼ ... mean radius mean texture mean perimeter mean area mean smoothness \
0      17.99      10.38      122.80      1001.0      0.11840
1      20.57      17.77      132.90      1326.0      0.08474
2      19.69      21.25      130.00      1203.0      0.10960
3      11.42      20.38       77.58       386.1      0.14250
4      20.29      14.34      135.10      1297.0      0.10030

      mean compactness mean concavity mean concave points mean symmetry \
0      0.27760      0.3001      0.14710      0.2419
1      0.07864      0.0869      0.07017      0.1812
2      0.15990      0.1974      0.12790      0.2069
3      0.28390      0.2414      0.10520      0.2597
4      0.13280      0.1980      0.10430      0.1809

      mean fractal dimension ... worst texture worst perimeter worst area \
0      0.07871 ...      17.33      184.60      2019.0
1      0.05667 ...      23.41      158.80      1956.0
2      0.05999 ...      25.53      152.50      1709.0
3      0.09744 ...      26.50       98.87       567.7
4      0.05883 ...      16.67      152.20      1575.0
```

```
      worst smoothness worst compactness worst concavity worst concave points \
0      0.1622      0.6656      0.7119      0.2654
1      0.1238      0.1866      0.2416      0.1860
2      0.1444      0.4245      0.4504      0.2430
3      0.2098      0.8663      0.6869      0.2575
4      0.1374      0.2050      0.4000      0.1625

      worst symmetry worst fractal dimension target
0      0.4601      0.11890      0
1      0.2750      0.08902      0
2      0.3613      0.08758      0
3      0.6638      0.17300      0
4      0.2364      0.07678      0
```

```
[5 rows x 31 columns]
```



```
df.isnull().sum()
```

...

0

mean radius

0

mean texture

0

mean perimeter

0

mean area

0

mean smoothness

0

mean compactness

0

mean concavity

0

mean concave points

0

mean symmetry

0

mean fractal dimension

0

radius error

0

texture error

0

perimeter error

0

area error	0
smoothness error	0
compactness error	0
concavity error	0
concave points error	0
symmetry error	0
fractal dimension error	0
worst radius	0
worst texture	0
worst perimeter	0
worst area	0
worst smoothness	0
worst compactness	0
worst concavity	0
worst concave points	0
worst symmetry	0

worst fractal dimension	0
target	0
dtype: int64	

```
[1] df.info()
```

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

#	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

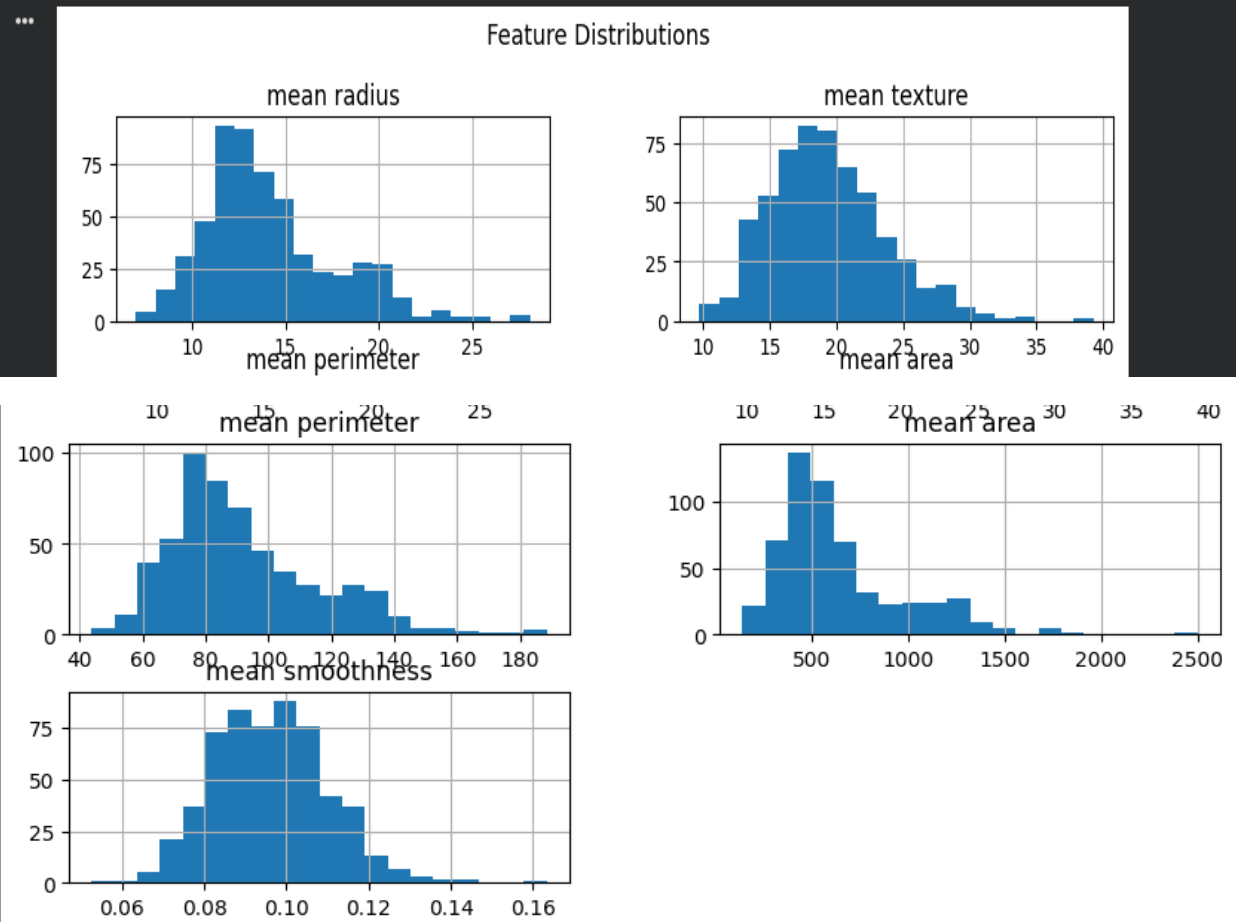
dtypes: float64(30), int64(1)  
memory usage: 137.9 KB

```
df.describe()
```

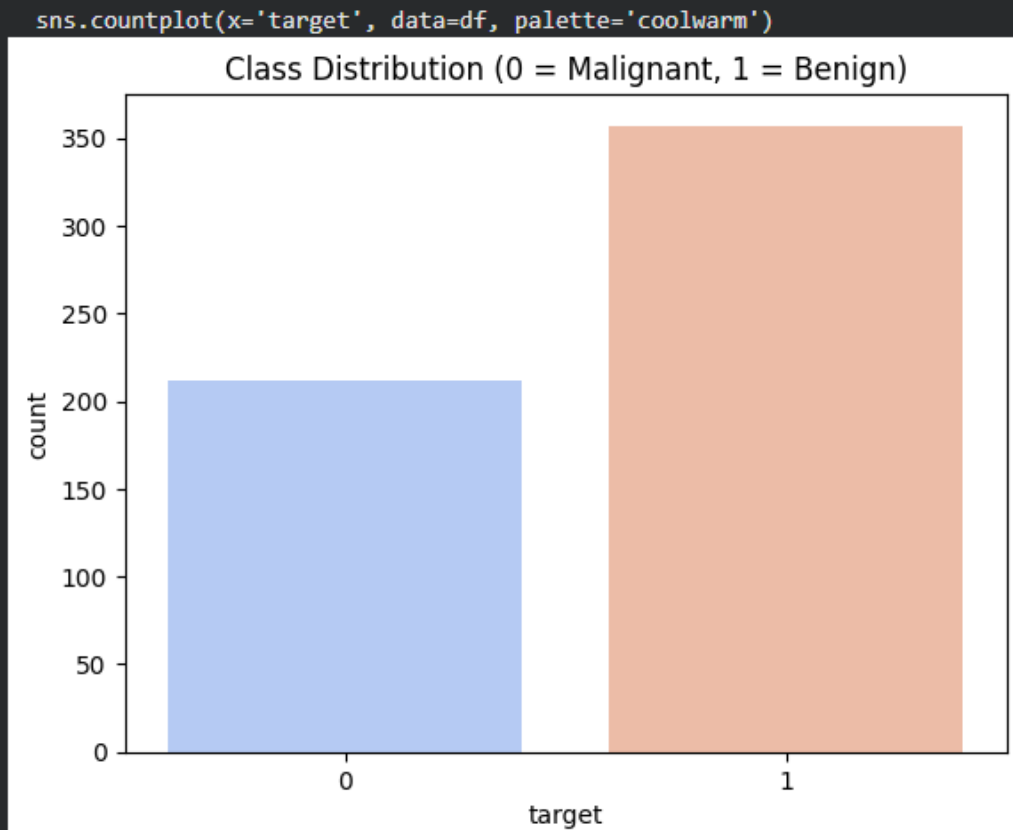
	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points	mean symmetry	mean fractal dimension	...	worst texture	worst perimeter
count	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	...	569.000000	569.000000
mean	14.127292	19.289649	91.969033	654.889104	0.096360	0.104341	0.088799	0.048919	0.181162	0.062798	...	25.677223	107.261213
std	3.524049	4.301036	24.298981	351.914129	0.014064	0.052813	0.079720	0.038803	0.027414	0.007060	...	6.146258	33.602542
min	6.981000	9.710000	43.790000	143.500000	0.052630	0.019380	0.000000	0.000000	0.106000	0.049960	...	12.020000	50.410000
25%	11.700000	16.170000	75.170000	420.300000	0.086370	0.064920	0.029560	0.020310	0.161900	0.057700	...	21.080000	84.110000
50%	13.370000	18.840000	86.240000	551.100000	0.095870	0.092630	0.061540	0.033500	0.179200	0.061540	...	25.410000	97.660000
75%	15.780000	21.800000	104.100000	782.700000	0.105300	0.130400	0.130700	0.074000	0.195700	0.066120	...	29.720000	125.400000
max	28.110000	39.280000	188.500000	2501.000000	0.163400	0.345400	0.426800	0.201200	0.304000	0.097440	...	49.540000	251.200000

8 rows x 31 columns

```
df.iloc[:, :5].hist(bins=20, figsize=(10, 6))
plt.suptitle("Feature Distributions")
plt.show()
```



```
▶ sns.countplot(x='target', data=df, palette='coolwarm')  
plt.title("Class Distribution (0 = Malignant, 1 = Benign)")  
plt.show()
```



#### 4. Data Preprocessing

- Split features and labels.
- Standardize features using StandardScaler.
- Train-test split.

```
[ ]  
X = df.drop('target', axis=1)  
y = df['target']  
  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42, stratify=y)
```

```

1 scaler = StandardScaler()
  X_train_scaled = scaler.fit_transform(X_train)
  X_test_scaled = scaler.transform(X_test)

```

## 5. Model Implementation

- Use `sklearn.linear_model.LogisticRegression`.
- Train the model on the training data.
- Predict on the test data.

```

model = LogisticRegression(max_iter=1000, random_state=42)
model.fit(X_train_scaled, y_train)

# Predict on test data
y_pred = model.predict(X_test_scaled)

```

## 6. Evaluation

- Accuracy score.
- Confusion matrix.
- Precision, Recall, F1-score

```

acc = accuracy_score(y_test, y_pred)
print(f"Accuracy: {acc*100:.2f}%")

```

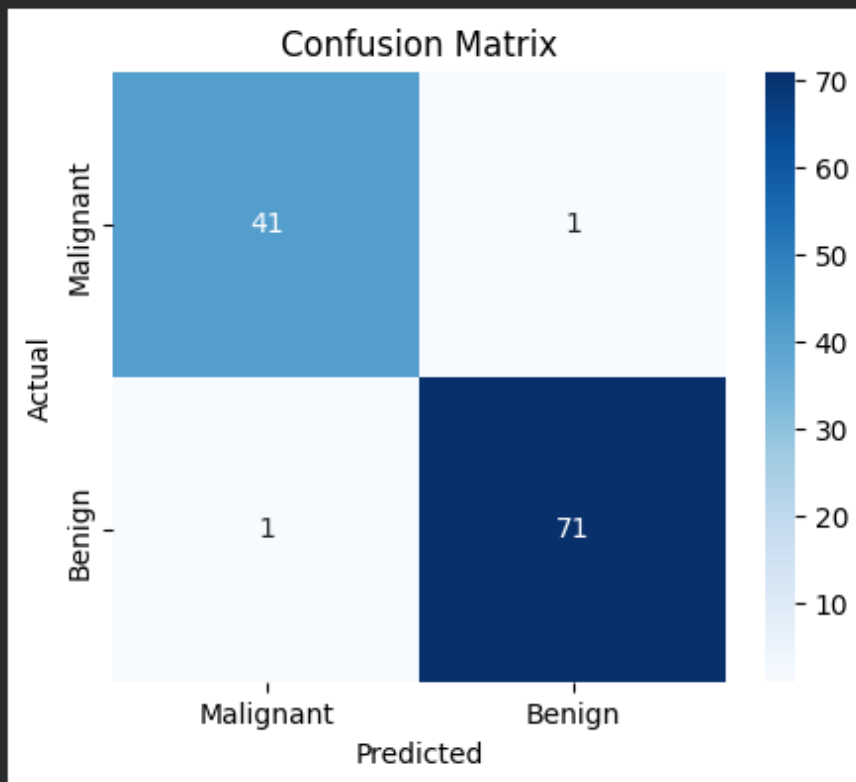
Accuracy: 98.25%

```

cm = confusion_matrix(y_test, y_pred)
plt.figure(figsize=(5,4))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=['Malignant', 'Benign'], yticklabels=['Malignant', 'Benign'])
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.title("Confusion Matrix")
plt.show()

```





```
print("\n ♦ Classification Report:")  
print(classification_report(y_test, y_pred, target_names=['Malignant', 'Benign']))
```

```
 ♦ Classification Report:  
              precision    recall  f1-score   support  
  
 Malignant      0.98      0.98      0.98        42  
 Benign         0.99      0.99      0.99        72  
  
 accuracy              0.98        114  
 macro avg           0.98      0.98      0.98        114  
 weighted avg        0.98      0.98      0.98        114
```