

BREAST CANCER CLASSIFICATION

- Algorithm: K-Nearest Neighbors (KNN), Random Forest Classifier, Linear Regression, Decision Tree Classifier, Logistic Regression
- Description: Classify breast cancer tumors as malignant or benign using features extracted from mammograms.
- For dataset-[here](#)

```
In [156]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
import warnings
warnings.filterwarnings('ignore')
```

```
In [157]: # Load the data
df = pd.read_csv('data.csv')
df
```

```
Out[157]:
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean
0	842302	M	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.30010
1	842517	M	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.08690
2	84300903	M	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.19740
3	84348301	M	11.42	20.38	77.58	386.1	0.14250	0.28390	0.24140
4	84358402	M	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.19800
...
564	926424	M	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390
565	926682	M	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400
566	926954	M	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251
567	927241	M	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140
568	92751	B	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000

569 rows × 33 columns

```
In [158]: df.head()
```

```
Out[158]:
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean
0	842302	M	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3001
1	842517	M	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0869
2	84300903	M	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.1974
3	84348301	M	11.42	20.38	77.58	386.1	0.14250	0.28390	0.2414
4	84358402	M	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1980

5 rows × 33 columns

```
In [159]: df.tail()
```

Out[159]:

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	
564	926424	M	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390	
565	926682	M	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400	
566	926954	M	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251	
567	927241	M	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140	
568	92751	B	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000	

5 rows × 33 columns

In [160]:

df.isnull().sum()

Out[160]:

id	0
diagnosis	0
radius_mean	0
texture_mean	0
perimeter_mean	0
area_mean	0
smoothness_mean	0
compactness_mean	0
concavity_mean	0
concave points_mean	0
symmetry_mean	0
fractal_dimension_mean	0
radius_se	0
texture_se	0
perimeter_se	0
area_se	0
smoothness_se	0
compactness_se	0
concavity_se	0
concave points_se	0
symmetry_se	0
fractal_dimension_se	0
radius_worst	0
texture_worst	0
perimeter_worst	0
area_worst	0
smoothness_worst	0
compactness_worst	0
concavity_worst	0
concave points_worst	0
symmetry_worst	0
fractal_dimension_worst	0
Unnamed: 32	569
dtype:	int64

In [161]:

df.info()

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 33 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   id                                     569 non-null    int64
1   diagnosis                             569 non-null    object
2   radius_mean                           569 non-null    float64
3   texture_mean                           569 non-null    float64
4   perimeter_mean                         569 non-null    float64
5   area_mean                             569 non-null    float64
6   smoothness_mean                       569 non-null    float64
7   compactness_mean                      569 non-null    float64
8   concavity_mean                        569 non-null    float64
9   concave points_mean                   569 non-null    float64
10  symmetry_mean                         569 non-null    float64
11  fractal_dimension_mean                569 non-null    float64
12  radius_se                             569 non-null    float64
13  texture_se                             569 non-null    float64
14  perimeter_se                           569 non-null    float64
15  area_se                               569 non-null    float64
16  smoothness_se                         569 non-null    float64
17  compactness_se                        569 non-null    float64
18  concavity_se                          569 non-null    float64
19  concave points_se                     569 non-null    float64
20  symmetry_se                           569 non-null    float64
21  fractal_dimension_se                  569 non-null    float64
22  radius_worst                          569 non-null    float64
23  texture_worst                         569 non-null    float64
24  perimeter_worst                       569 non-null    float64
25  area_worst                            569 non-null    float64
26  smoothness_worst                     569 non-null    float64
27  compactness_worst                     569 non-null    float64
28  concavity_worst                       569 non-null    float64
29  concave points_worst                  569 non-null    float64
30  symmetry_worst                        569 non-null    float64
31  fractal_dimension_worst                569 non-null    float64
32  Unnamed: 32                           0 non-null      float64
dtypes: float64(31), int64(1), object(1)
memory usage: 146.8+ KB

```

In [162]: df.describe()

```

Out[162]:

```

	id	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	po
count	5.690000e+02	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	569.000000	5
mean	3.037183e+07	14.127292	19.289649	91.969033	654.889104	0.096360	0.104341	0.088799	
std	1.250206e+08	3.524049	4.301036	24.298981	351.914129	0.014064	0.052813	0.079720	
min	8.670000e+03	6.981000	9.710000	43.790000	143.500000	0.052630	0.019380	0.000000	
25%	8.692180e+05	11.700000	16.170000	75.170000	420.300000	0.086370	0.064920	0.029560	
50%	9.060240e+05	13.370000	18.840000	86.240000	551.100000	0.095870	0.092630	0.061540	
75%	8.813129e+06	15.780000	21.800000	104.100000	782.700000	0.105300	0.130400	0.130700	
max	9.113205e+08	28.110000	39.280000	188.500000	2501.000000	0.163400	0.345400	0.426800	

8 rows × 32 columns

```

In [163]: print("Number of rows",df.shape[0])
          print("Number of columns",df.shape[1])

```

```

Number of rows 569
Number of columns 33

```

```

In [164]: print(df.duplicated().sum())

```

```

0

```

```

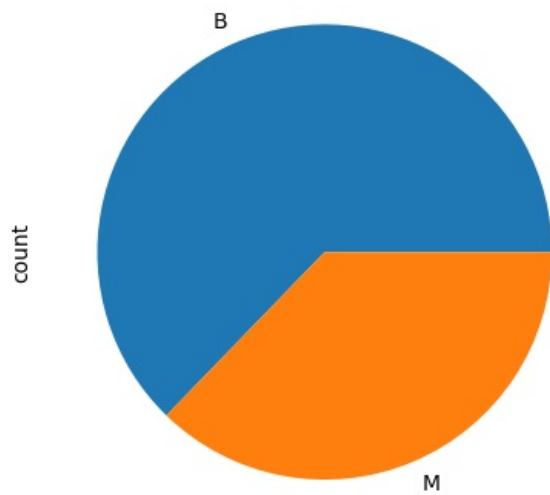
In [165]: #visualization
          df['diagnosis'].value_counts().plot(kind='pie')

```

```

Out[165]: <Axes: ylabel='count'>

```



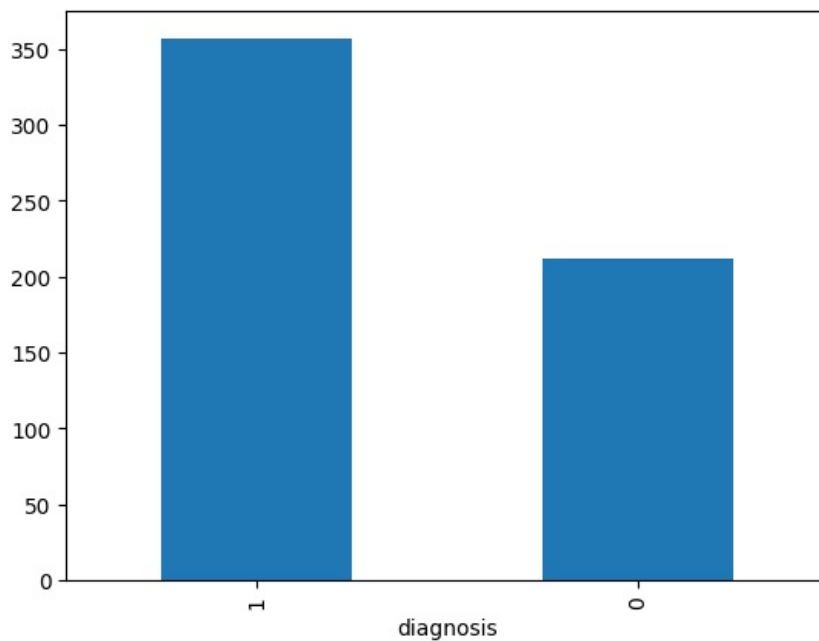
```
In [166.. # Data Preprocessing
df['diagnosis'] = df['diagnosis'].map({"B": 1, "M": 0})
```

```
In [167.. df['diagnosis'].value_counts()
```

```
Out[167]: diagnosis
1      357
0      212
Name: count, dtype: int64
```

```
In [168.. #visualization using barplot
df['diagnosis'].value_counts().plot(kind='bar')
```

```
Out[168]: <Axes: xlabel='diagnosis'>
```



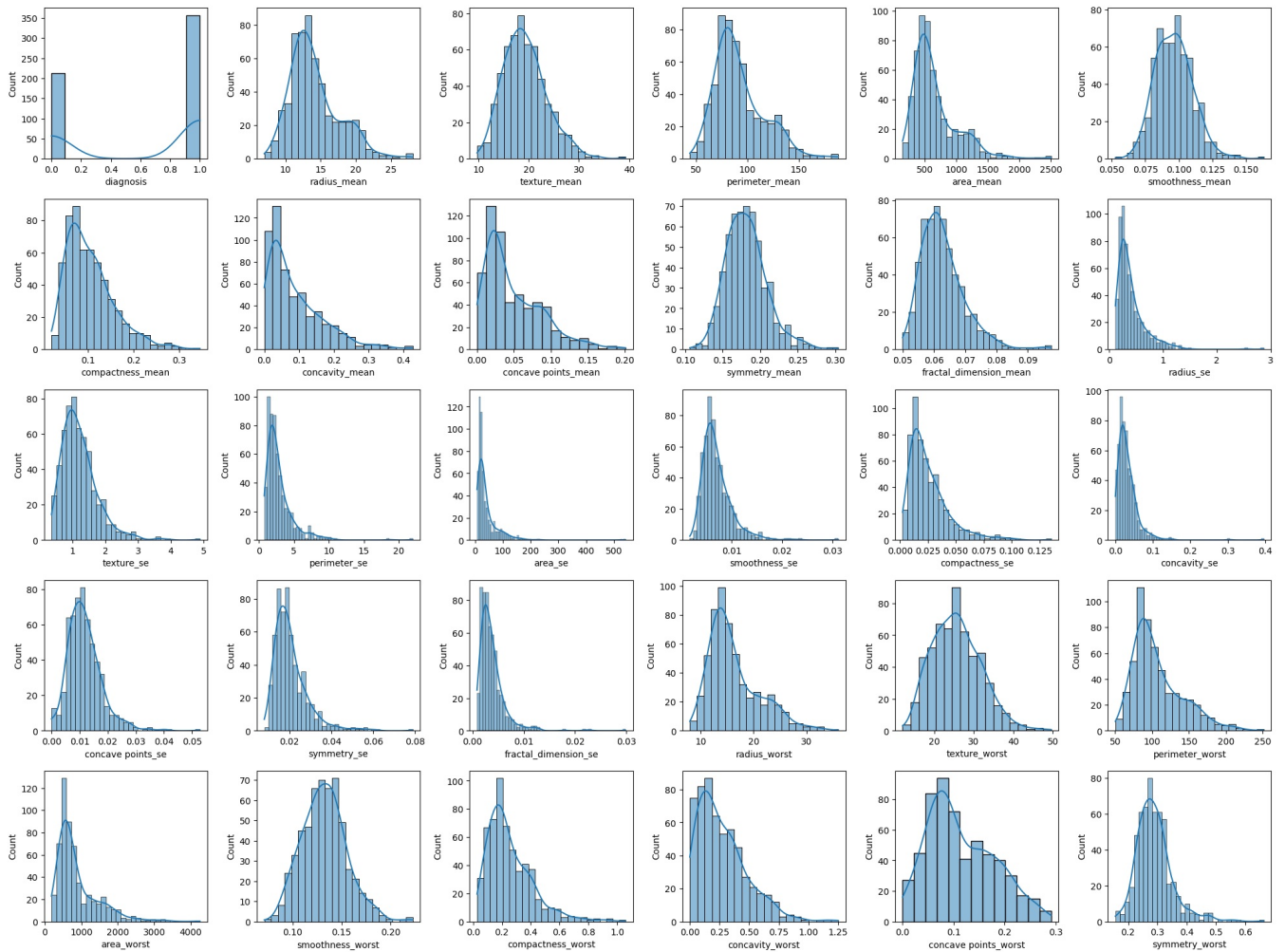
```
In [169.. #drop unnecessary feature
df.drop(['id', 'Unnamed: 32'], axis=1, inplace=True)
```

```
In [170.. # Visualize the distribution of features
plt.figure(figsize=(20, 15))
plotnumber = 1
for column in df.columns:
    if plotnumber <= 30:
```

```

ax = plt.subplot(5, 6, plotnumber)
sns.histplot(df[column], kde=True)
plt.xlabel(column)
plotnumber += 1
plt.tight_layout()
plt.show()

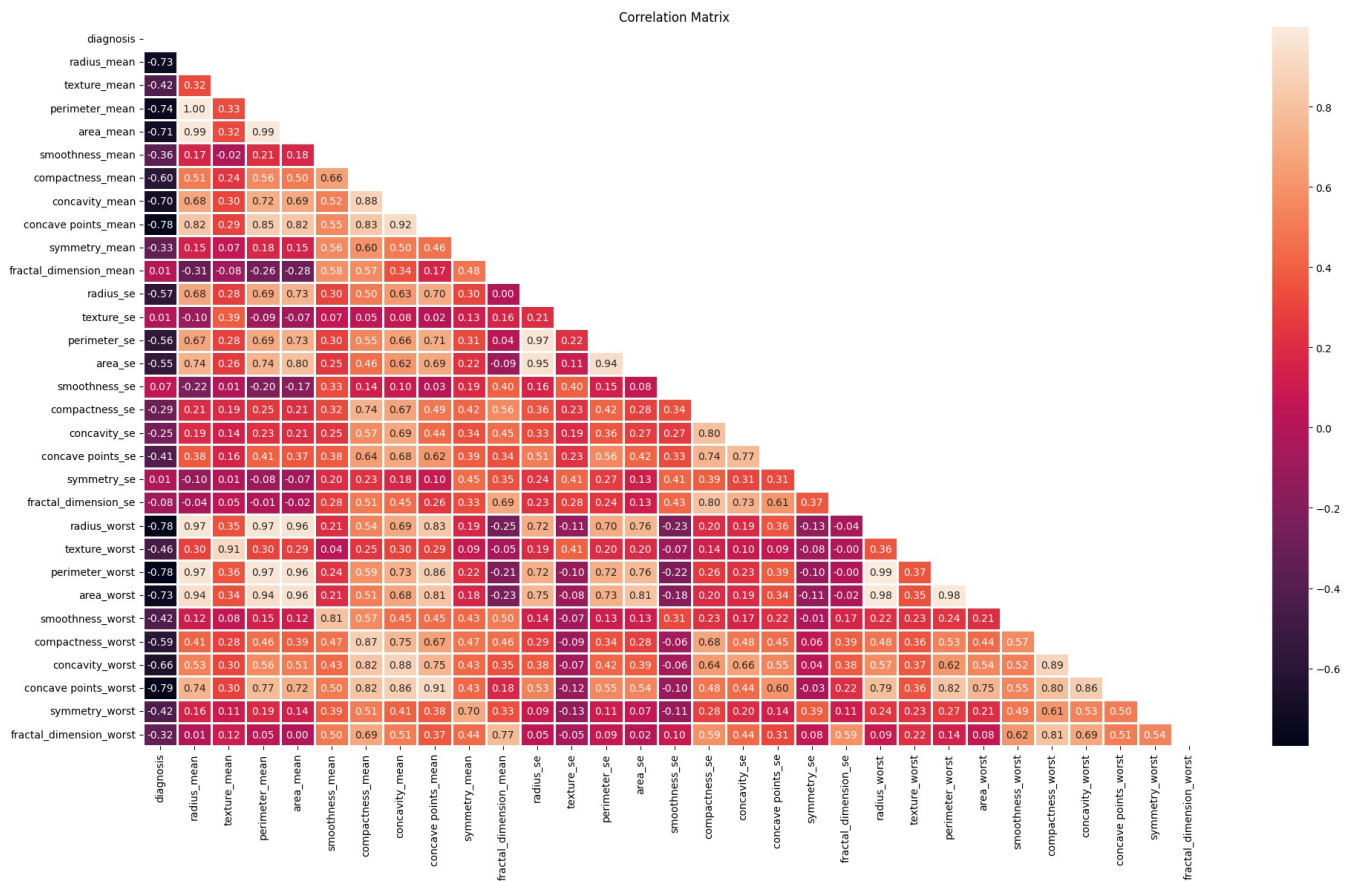
```



```

In [171]: # Heatmap of correlations
plt.figure(figsize=(20, 12))
corr = df.corr()
mask = np.triu(np.ones_like(corr, dtype=bool))
sns.heatmap(corr, mask=mask, linewidths=1, annot=True, fmt=".2f")
plt.title("Correlation Matrix")
plt.tight_layout()
plt.show()

```



```
In [172.. # Drop highly correlated features
corr_matrix = df.corr().abs()
mask = np.triu(np.ones_like(corr_matrix, dtype=bool))
tri_df = corr_matrix.mask(mask)
to_drop = [x for x in tri_df.columns if any(tri_df[x] > 0.92)]
df.drop(to_drop, axis=1, inplace=True)
print(f"Number of features after dropping correlated ones: {df.shape[1]}")
```

Number of features after dropping correlated ones: 23

```
In [173.. # Prepare data for training
X = df.drop('diagnosis', axis=1)
y = df['diagnosis']
```

```
In [174.. X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
scaler = StandardScaler()
X_train = scaler.fit_transform(X_train)
X_test = scaler.transform(X_test)
```

```
In [175.. # Define models
models = [
    ("Logistic Regression", LogisticRegression(max_iter=200)),
    ("Random Forest", RandomForestClassifier(random_state=42)),
    ("K-Nearest Neighbors", KNeighborsClassifier()),
    ("Decision Tree", DecisionTreeClassifier(random_state=42)),
]
```

```
In [176.. # Train and evaluate models, storing results
results = {}
for name, model in models:
    model.fit(X_train, y_train)
    predictions = model.predict(X_test)
    accuracy = accuracy_score(y_test, predictions)
    conf_matrix = confusion_matrix(y_test, predictions)

    results[name] = {
        "Accuracy": accuracy,
        "Confusion Matrix": conf_matrix
    }

# Print results for each model
print(f"{name} Accuracy: {accuracy:.4f}")
print(classification_report(y_test, predictions, target_names=["Malignant", "Benign"]))
print("-" * 50)
```

Logistic Regression Accuracy: 0.9737				
	precision	recall	f1-score	support
Malignant	0.98	0.95	0.96	43
Benign	0.97	0.99	0.98	71
accuracy			0.97	114
macro avg	0.97	0.97	0.97	114
weighted avg	0.97	0.97	0.97	114

Random Forest Accuracy: 0.9649				
	precision	recall	f1-score	support
Malignant	0.98	0.93	0.95	43
Benign	0.96	0.99	0.97	71
accuracy			0.96	114
macro avg	0.97	0.96	0.96	114
weighted avg	0.97	0.96	0.96	114

K-Nearest Neighbors Accuracy: 0.9386				
	precision	recall	f1-score	support
Malignant	0.93	0.91	0.92	43
Benign	0.94	0.96	0.95	71
accuracy			0.94	114
macro avg	0.94	0.93	0.93	114
weighted avg	0.94	0.94	0.94	114

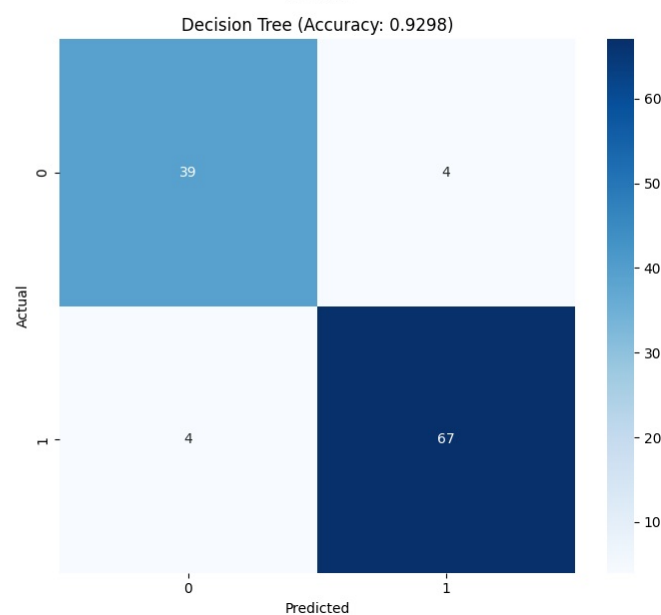
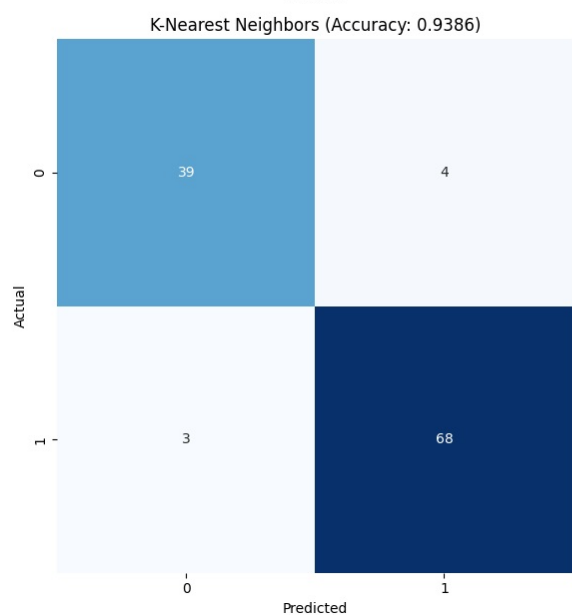
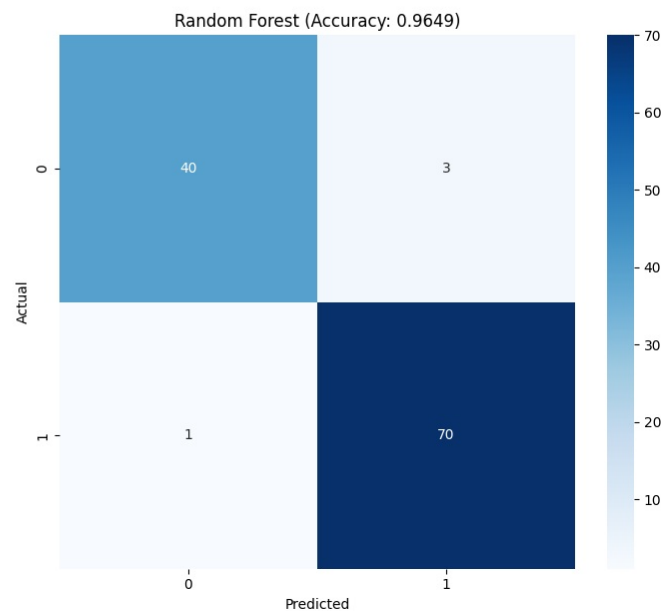
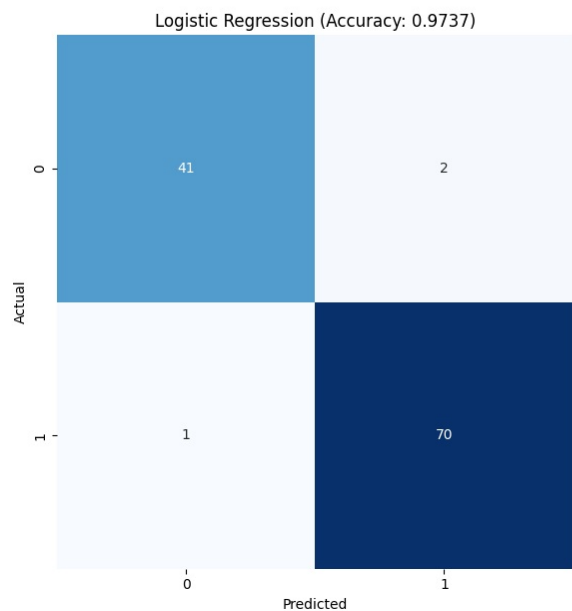
Decision Tree Accuracy: 0.9298				
	precision	recall	f1-score	support
Malignant	0.91	0.91	0.91	43
Benign	0.94	0.94	0.94	71
accuracy			0.93	114
macro avg	0.93	0.93	0.93	114
weighted avg	0.93	0.93	0.93	114

```
In [177]: # Plot the confusion matrices
fig, axes = plt.subplots(2, 2, figsize=(14, 14))
fig.suptitle('Confusion Matrices of Different Models', fontsize=20)

for (name, result), ax in zip(results.items(), axes.flatten()):
    sns.heatmap(result['Confusion Matrix'], annot=True, fmt='d', cmap='Blues', ax=ax)
    ax.set_title(f'{name} (Accuracy: {result["Accuracy"]:.4f})')
    ax.set_xlabel('Predicted')
    ax.set_ylabel('Actual')

plt.tight_layout(rect=[0, 0.03, 1, 0.95])
plt.show()
```

Confusion Matrices of Different Models



In []:

In []:

In []:

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