

CODE

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import classification_report, accuracy_score
from sklearn.preprocessing import LabelEncoder
import matplotlib.pyplot as plt
import seaborn as sns

# Step 1: Load Data
try:
    data = pd.read_csv('breast_cancer_data.csv')
    print("Data loaded successfully")
except Exception as e:
    print(f"Error loading data: {e}")

# View the first few rows of the dataset
print("First few rows of the dataset:")
print(data.head())

# Step 2: Define Features and Target
try:
    X = data[['radius_mean', 'texture_mean', 'perimeter_mean', 'area_mean',
              'smoothness_mean', 'compactness_mean', 'concavity_mean',
              'concave_points_mean', 'symmetry_mean', 'fractal_dimension_mean']]
    y = data['diagnosis']
    print("Features and target variable defined successfully")
except KeyError as e:
    print(f"Error defining features and target: {e}")

# Encode the target variable
try:
    label_encoder = LabelEncoder()
    y_encoded = label_encoder.fit_transform(y) # Convert 'M' and 'B' to 1 and 0
    print("Target variable encoded successfully")
except Exception as e:
    print(f"Error encoding target variable: {e}")
```

Step 3: Split the Data into Training and Testing Sets

try:

```
X_train, X_test, y_train, y_test = train_test_split(X, y_encoded, test_size=0.2,
random_state=0)
```

```
print("Data split into training and testing sets successfully")
```

except Exception as e:

```
print(f"Error splitting data: {e}")
```

Define and Train the Model

try:

```
model = RandomForestClassifier(n_estimators=100, random_state=0)
```

```
model.fit(X_train, y_train)
```

```
print("Model trained successfully")
```

except Exception as e:

```
print(f"Error training the model: {e}")
```

Make Predictions

try:

```
y_pred = model.predict(X_test)
```

```
print("Predictions made successfully")
```

except Exception as e:

```
print(f"Error making predictions: {e}")
```

Evaluate the Model

try:

```
print("Accuracy:", accuracy_score(y_test, y_pred))
```

```
print("Classification Report:\n", classification_report(y_test, y_pred))
```

except Exception as e:

```
print(f"Error evaluating the model: {e}")
```

Feature Importances

try:

```
importances = model.feature_importances_
```

```
features = X.columns
```

```
feature_importance_df = pd.DataFrame({
```

```
    'Feature': features,
```

```

        'Importance': importances
    }).sort_values(by='Importance', ascending=False)
    print("Feature importances calculated successfully")
except Exception as e:
    print(f"Error calculating feature importances: {e}")
import matplotlib.pyplot as plt
import seaborn as sns

# Assuming 'y_encoded' contains the encoded target variable (0: Benign, 1: Malignant)

# Plot the distribution of diagnosis (Malignant and Benign)
plt.figure(figsize=(8, 6))
sns.countplot(x=y_encoded, palette="coolwarm")
plt.title('Distribution of Breast Cancer Diagnosis')
plt.xlabel('Diagnosis (0: Benign, 1: Malignant)')
plt.ylabel('Count')
plt.show()

# Step 7: Identify and Print Individuals Affected by Cancer
try:
    malignant_indices = X_test.index[y_pred == 1]
    if not malignant_indices.empty:
        malignant_cases = data.loc[malignant_indices]
        print("Individuals affected by cancer (predicted malignant cases):")
        print(malignant_cases)
    else:
        print("No individuals predicted to be affected by cancer in the test set.")
except Exception as e:
    print(f"Error identifying malignant cases: {e}")

```

OUTPUT

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	\
0	842302	M	17.99	10.38	122.80	1001.0	
1	842517	M	20.57	17.77	132.90	1326.0	
2	84300903	M	19.69	21.25	130.00	1203.0	
3	84348301	M	11.42	20.38	77.58	386.1	
4	84358402	M	20.29	14.34	135.10	1297.0	

	smoothness_mean	compactness_mean	concavity_mean	concave_points_mean	\
0	0.11840	0.27760	0.3001	0.14710	
1	0.08474	0.07864	0.0869	0.07017	
2	0.10960	0.15990	0.1974	0.12790	
3	0.14250	0.28390	0.2414	0.10520	
4	0.10030	0.13280	0.1980	0.10430	

	...	radius_worst	texture_worst	perimeter_worst	area_worst	\
0	...	25.38	17.33	184.60	2019.0	
1	...	24.99	23.41	158.80	1956.0	
2	...	23.57	25.53	152.50	1709.0	
3	...	14.91	26.50	98.87	567.7	
4	...	22.54	16.67	152.20	1575.0	

	smoothness_worst	compactness_worst	concavity_worst	concave_points_worst	
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	

	symmetry_worst	fractal_dimension_worst	
0	0.4601	0.11890	
1	0.2750	0.08902	
2	0.3613	0.08758	
3	0.6638	0.17300	
4	0.2364	0.07678	

[5 rows x 32 columns]
Accuracy: 0.9298245614035088
Classification Report:

	precision	recall	f1-score	support
0	0.95	0.93	0.94	67
1	0.90	0.94	0.92	47

accuracy			0.93	114
macro avg	0.93	0.93	0.93	114
weighted avg	0.93	0.93	0.93	114

AFFECTED PEOPLE

Individuals affected by cancer (predicted malignant cases):

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	\	area_mean	smoothness_mean	compactness_mean	concavity_mean	\
512	915691	M	13.40	20.52	88.64		556.7	0.11060	0.14690	0.14450	
421	906564	B	14.69	13.98	98.22		656.1	0.10310	0.18360	0.14500	
157	8711216	B	16.84	19.46	108.40		880.2	0.07445	0.07223	0.05150	
89	861598	B	14.64	15.24	95.77		651.9	0.11320	0.13390	0.09966	
172	87164	M	15.46	11.89	102.50		736.9	0.12570	0.15550	0.20320	
233	88206102	M	20.51	27.81	134.40		1319.0	0.09159	0.10740	0.15540	
389	90312	M	19.55	23.21	128.90		1174.0	0.10100	0.13180	0.18560	
250	884948	M	20.94	23.56	138.90		1364.0	0.10070	0.16060	0.27120	
283	8912280	M	16.24	18.77	108.80		805.1	0.10660	0.18020	0.19480	
372	9012795	M	21.37	15.10	141.30		1386.0	0.10010	0.15150	0.19320	
14	84667401	M	13.73	22.61	93.60		578.3	0.11310	0.22930	0.21280	
337	897630	M	18.77	21.43	122.90		1092.0	0.09116	0.14020	0.10600	
1	842517	M	20.57	17.77	132.90		1326.0	0.08474	0.07864	0.08690	
132	86730502	M	16.16	21.54	106.20		809.8	0.10080	0.12840	0.10430	
64	85922302	M	12.68	23.84	82.69		499.0	0.11220	0.12620	0.11280	
127	866203	M	19.00	18.91	123.40		1138.0	0.08217	0.08028	0.09271	
353	9010018	M	15.08	25.74	98.00		716.6	0.10240	0.09769	0.12350	
414	905680	M	15.13	29.81	96.71		719.5	0.08320	0.04605	0.04686	
10	845636	M	16.02	23.24	102.70		797.8	0.08206	0.06669	0.03299	
564	926424	M	21.56	22.39	142.00		1479.0	0.11100	0.11590	0.24390	
15	84799002	M	14.54	27.54	96.73		658.8	0.11390	0.15950	0.16390	
12	846226	M	19.17	24.80	132.40		1123.0	0.09740	0.24580	0.20650	
194	87556202	M	14.86	23.21	100.40		671.4	0.10440	0.19800	0.16970	
134	867739	M	18.45	21.91	120.20		1075.0	0.09430	0.09709	0.11530	
272	8910988	M	21.75	20.99	147.30		1491.0	0.09401	0.19610	0.21950	
196	875938	M	13.77	22.29	90.63		588.9	0.12000	0.12670	0.13850	
75	8610404	M	16.07	19.65	104.10		817.7	0.09168	0.08424	0.09769	
468	9113538	M	17.60	23.33	119.00		980.5	0.09289	0.20040	0.21360	
108	86355	M	22.27	19.67	152.80		1509.0	0.13260	0.27680	0.42640	
239	88330202	M	17.46	39.28	113.40		920.6	0.09812	0.12980	0.14170	
210	881046502	M	20.58	22.14	134.70		1290.0	0.09090	0.13480	0.16400	
							798.8	0.11700	0.20220	0.17220	
							664.7	0.08682	0.06636	0.08390	

