

```
# Import Libraries
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

```
# Load the dataset
df=pd.read_csv('/content/breast-cancer.csv')
```

```
# Display first few rows
df.head()
```



	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothn
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	

5 rows × 32 columns

```
# Display basic information
df.info()
```



#	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
dtypes: float64(30), int64(1), object(1)
memory usage: 142.4+ KB

```

1. Load and prepare a dataset for binary classification

```

# Drop the 'id' column
df.drop('id',axis=1,inplace=True)

# Encode 'diagnosis' as binary
df['diagnosis']=df['diagnosis'].map({'M':1,'B':0})

```

```

# Split data into features and target
X=df.drop(['diagnosis'],axis=1)
y=df['diagnosis']

```

```

# Normalize the features
from sklearn.preprocessing import StandardScaler

```

```
ss=StandardScaler()
```

```
X=ss.fit_transform(X)
```

```
# Train-test split
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test= train_test_split(X,y,test_size=0.2,random_state=42)
```

```
X_train.shape, X_test.shape, y_train.shape, y_test.shape
```

→ ((455, 31), (114, 31), (455,), (114,))

2. Train an SVM with linear and RBF kernel.

```
from sklearn.svm import SVC
```

```
# Train SVM with linear kernel
svm_linear = SVC(kernel='linear', random_state=42)
svm_linear.fit(X_train, y_train)
y_pred_linear = svm_linear.predict(X_test)

# Train SVM with RBF kernel
svm_rbf = SVC(kernel='rbf', random_state=42)
svm_rbf.fit(X_train, y_train)
y_pred_rbf = svm_rbf.predict(X_test)

# Evaluate both models
from sklearn.metrics import accuracy_score, classification_report, confusion_matrix

linear_results = {
    "Accuracy": accuracy_score(y_test, y_pred_linear),
    "Classification Report": classification_report(y_test, y_pred_linear, output_dict=True),
    "Confusion Matrix": confusion_matrix(y_test, y_pred_linear)
}
```

```
linear_results
```

```
→ {'Accuracy': 1.0,
    'Classification Report': {'0': {'precision': 1.0,
        'recall': 1.0,
        'f1-score': 1.0,
        'support': 71.0},
      '1': {'precision': 1.0, 'recall': 1.0, 'f1-score': 1.0, 'support': 43.0},
      'accuracy': 1.0,
      'macro avg': {'precision': 1.0,
        'recall': 1.0,
        'f1-score': 1.0,
        'support': 114.0},
      'weighted avg': {'precision': 1.0,
        'recall': 1.0,
        'f1-score': 1.0,
        'support': 114.0}},
    'Confusion Matrix': array([[71,  0],
       [ 0, 43]])}
```

```
rbf_results = {
    "Accuracy": accuracy_score(y_test, y_pred_rbf),
    "Classification Report": classification_report(y_test, y_pred_rbf, output_dict=True),
    "Confusion Matrix": confusion_matrix(y_test, y_pred_rbf)
}
```

```
rbf_results
```

```
→ {'Accuracy': 1.0,
    'Classification Report': {'0': {'precision': 1.0,
        'recall': 1.0,
        'f1-score': 1.0,
```

```
'support': 71.0},
'1': {'precision': 1.0, 'recall': 1.0, 'f1-score': 1.0, 'support': 43.0},
'accuracy': 1.0,
'macro avg': {'precision': 1.0,
'recall': 1.0,
'f1-score': 1.0,
'support': 114.0},
'weighted avg': {'precision': 1.0,
'recall': 1.0,
'f1-score': 1.0,
'support': 114.0}},
'Confusion Matrix': array([[71,  0],
[ 0, 43]])}
```

3. Visualize decision boundary using 2D data.

```
# Select two features for 2D visualization
feature1 = 'radius_mean'
feature2 = 'texture_mean'
```

```
# Extract corresponding columns and scale them
X_2d = df[[feature1, feature2]]
X_2d_scaled = ss.fit_transform(X_2d)
```

```
# Re-split for 2D visualization
X_train_2d, X_test_2d, y_train_2d, y_test_2d = train_test_split(X_2d_scaled, y, test_size
```

```
# Fit both SVMs on 2D data
svm_linear_2d = SVC(kernel='linear')
svm_linear_2d.fit(X_train_2d, y_train_2d)
```

```
SVC(kernel='linear')
```

```
svm_rbf_2d = SVC(kernel='rbf')
svm_rbf_2d.fit(X_train_2d, y_train_2d)
```

```
SVC()
```

```
# Function to plot decision boundary
def plot_decision_boundary(clf, X, y, title):
    h = .02 # step size in the mesh
    x_min, x_max = X[:, 0].min() - 1, X[:, 0].max() + 1
    y_min, y_max = X[:, 1].min() - 1, X[:, 1].max() + 1
    xx, yy = np.meshgrid(np.arange(x_min, x_max, h),
                         np.arange(y_min, y_max, h))
```

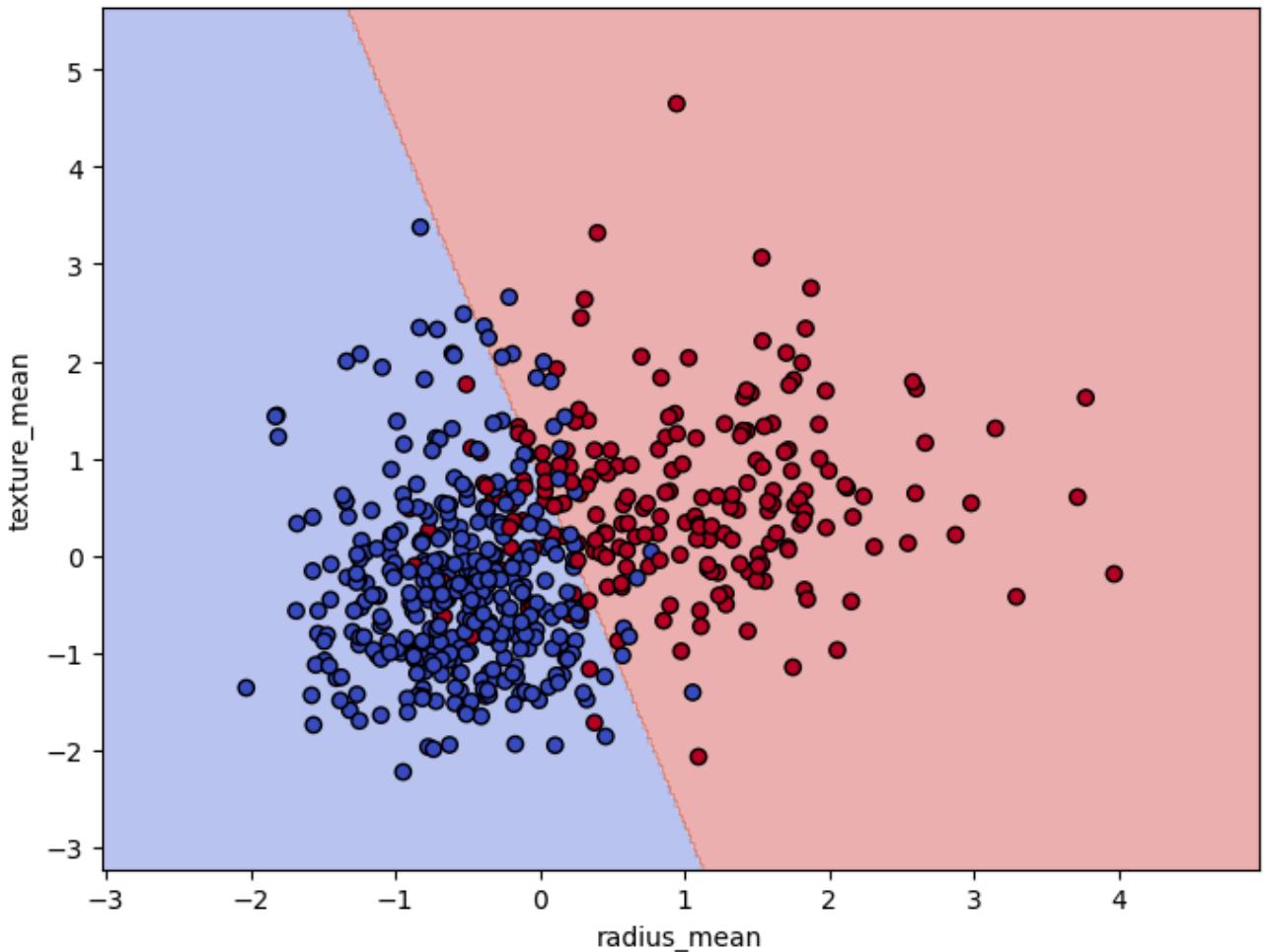
```
Z = clf.predict(np.c_[xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)

plt.figure(figsize=(8, 6))
plt.contourf(xx, yy, Z, alpha=0.4, cmap=plt.cm.coolwarm)
plt.scatter(X[:, 0], X[:, 1], c=y, cmap=plt.cm.coolwarm, edgecolors='k')
plt.xlabel(feature1)
plt.ylabel(feature2)
plt.title(title)
plt.show()
```

```
# Plot decision boundaries
plot_decision_boundary(svm_linear_2d, X_2d_scaled, y, "SVM with Linear Kernel (2D)")
```



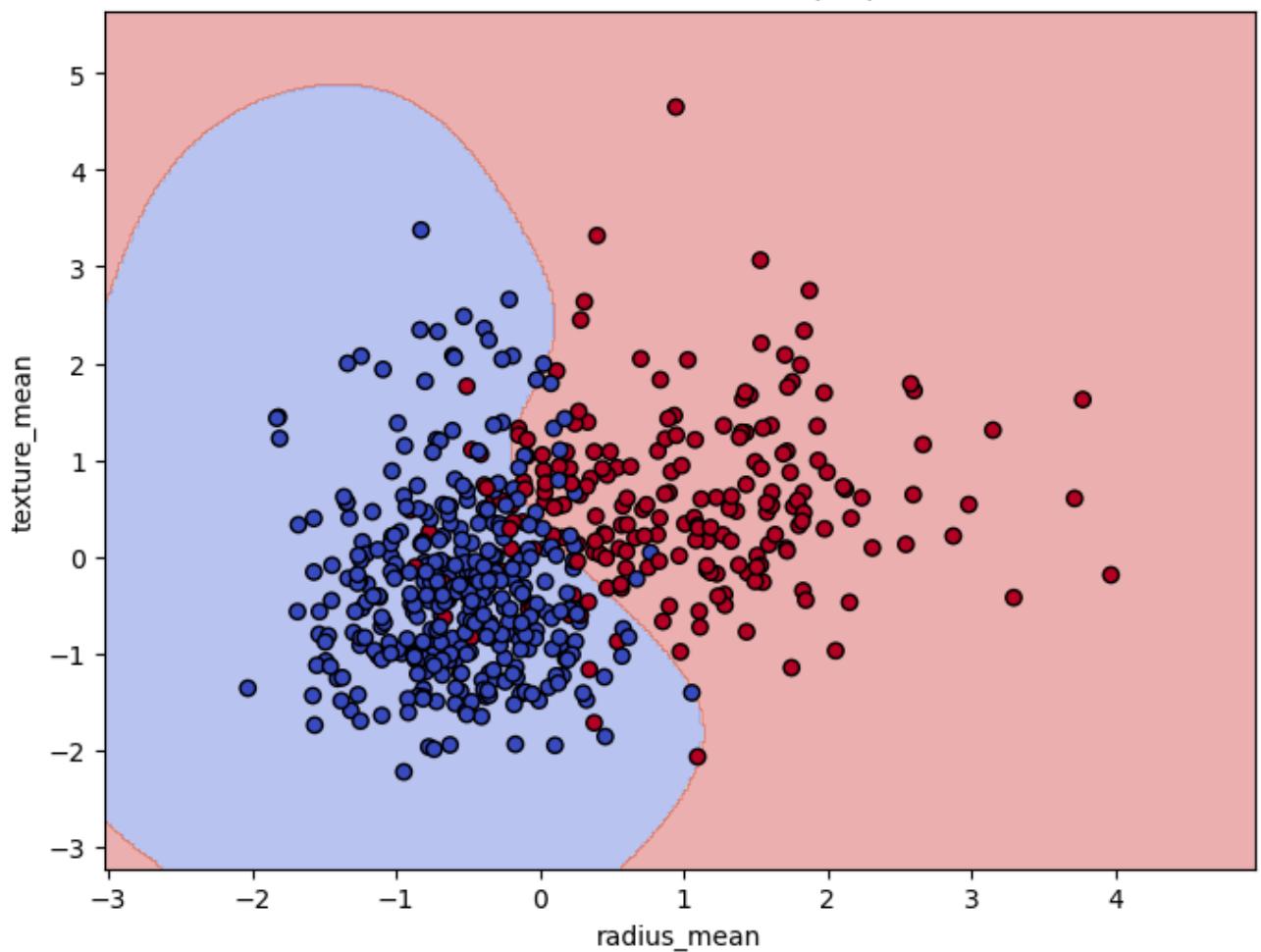
SVM with Linear Kernel (2D)



```
plot_decision_boundary(svm_rbf_2d, X_2d_scaled, y, "SVM with RBF Kernel (2D)")
```



SVM with RBF Kernel (2D)



4.Tune hyperparameters like C and gamma

```
from sklearn.model_selection import GridSearchCV

# Define parameter grid for SVM with RBF kernel
param_grid = {
    'C': [0.1, 1, 10, 100],
    'gamma': [0.01, 0.1, 1, 10],
    'kernel': ['rbf']
}

# Perform grid search with cross-validation
grid_search = GridSearchCV(SVC(), param_grid, cv=5, scoring='accuracy')
grid_search.fit(X_train, y_train)
```



```
# Print best parameters and accuracy
print("Best Parameters:", grid_search.best_params_)
```

→ Best Parameters: {'C': 10, 'gamma': 0.01, 'kernel': 'rbf'}

```
print("Best CV Accuracy:", grid_search.best_score_)
```

→ Best CV Accuracy: 1.0

```
# Test set performance
best_model = grid_search.best_estimator_
test_accuracy = best_model.score(X_test, y_test)
print("Test Accuracy:", test_accuracy)
```

→ Test Accuracy: 1.0

5. Use cross-validation to evaluate performance

```
from sklearn.model_selection import cross_val_score
```

```
# Linear SVM
svm_linear = SVC(kernel='linear', C=1)
linear_scores = cross_val_score(svm_linear, X, y, cv=5, scoring='accuracy')
```

```
# Print results
print("Linear SVM CV Accuracy: %0.4f ± %0.4f" % (linear_scores.mean(), linear_scores.std()))
```

→ Linear SVM CV Accuracy: 1.0000 ± 0.0000

```
# RBF SVM
svm_rbf = SVC(kernel='rbf', C=1, gamma='scale') # 'scale' is default in recent sklearn
rbf_scores = cross_val_score(svm_rbf, X, y, cv=5, scoring='accuracy')
```

```
# Print results
print("RBF SVM CV Accuracy: %0.4f ± %0.4f" % (rbf_scores.mean(), rbf_scores.std()))
```

→ RBF SVM CV Accuracy: 0.9965 ± 0.0070

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