ml-ps4

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```
import numpy as np
import pandas as pd
from sklearn.ensemble import RandomForestClassifier, RandomForestRegressor
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, precision_score, recall_score,

of1_score, roc_auc_score, roc_curve, mean_squared_error, r2_score
import matplotlib.pyplot as plt
import seaborn as sns
```

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[3]: breast_cancer_url = "https://archive.ics.uci.edu/ml/machine-learning-databases/
    ⇔breast-cancer-wisconsin/wdbc.data"
    breast_cancer_columns = ['ID', 'Diagnosis'] + [f'feature_{i}' for i in range(1, __
    breast_cancer_df = pd.read_csv(breast_cancer_url, header=None,_
     →names=breast_cancer_columns)
    # Prepare the dataset
    X_cancer = breast_cancer_df.drop(['ID', 'Diagnosis'], axis=1)
    y_cancer = breast_cancer_df['Diagnosis'].apply(lambda x: 1 if x == 'M' else 0)
    # Split the dataset into training and testing sets
    X_train_cancer, X_test_cancer, y_train_cancer, y_test_cancer =_
     # Load the California Housing Dataset
    housing_url = "https://download.mlcc.google.com/mledu-datasets/
     ⇔california_housing_train.csv"
    housing_df = pd.read_csv(housing_url)
    # Prepare the dataset
    X_housing = housing_df.drop(['median_house_value'], axis=1)
    y_housing = housing_df['median_house_value']
```

```
# Split the dataset into training and testing sets
     X train housing, X test housing, y train housing, y test housing = ___
      cutrain_test_split(X_housing, y_housing, test_size=0.3, random_state=42)
[4]: # Initialize the models
     rf_classifier = RandomForestClassifier(random_state=42)
     rf_regressor = RandomForestRegressor(random_state=42)
[5]: # Train the models
     rf_classifier.fit(X_train_cancer, y_train_cancer)
     rf_regressor.fit(X_train_housing, y_train_housing)
     # Make predictions
     y_pred_cancer = rf_classifier.predict(X_test_cancer)
     y pred proba cancer = rf classifier.predict proba(X test cancer)[:, 1]
     y_pred_housing = rf_regressor.predict(X_test_housing)
[6]: # Classification Metrics
     accuracy = accuracy_score(y_test_cancer, y_pred_cancer)
     precision = precision_score(y_test_cancer, y_pred_cancer)
     recall = recall score(y test cancer, y pred cancer)
     f1 = f1_score(y_test_cancer, y_pred_cancer)
     roc_auc = roc_auc_score(y_test_cancer, y_pred_proba_cancer)
     # Regression Metrics
     mse = mean_squared_error(y_test_housing, y_pred_housing)
     r_squared = r2_score(y_test_housing, y_pred_housing)
     # Output metrics
     print("Classification Metrics")
     print(f"Accuracy: {accuracy}")
     print(f"Precision: {precision}")
     print(f"Recall: {recall}")
     print(f"F1 Score: {f1}")
     print(f"ROC AUC: {roc_auc}")
     print("\nRegression Metrics")
     print(f"Mean Squared Error: {mse}")
```

Classification Metrics

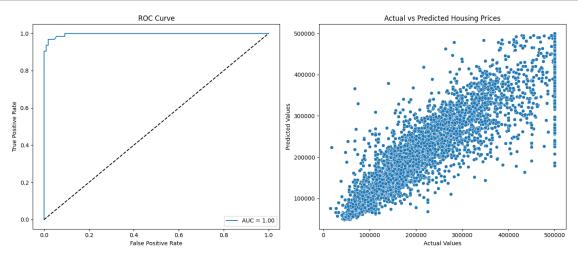
print(f"R-Squared: {r_squared}")

Regression Metrics

Mean Squared Error: 2560410737.09187

R-Squared: 0.8146160365959992

```
[7]: # Plot ROC curve for Classification Task
     fpr, tpr, _ = roc_curve(y_test_cancer, y_pred_proba_cancer)
     plt.figure(figsize=(14, 6))
     plt.subplot(1, 2, 1)
     plt.plot(fpr, tpr, label=f'AUC = {roc_auc:.2f}')
     plt.plot([0, 1], [0, 1], 'k--')
     plt.xlabel('False Positive Rate')
     plt.ylabel('True Positive Rate')
     plt.title('ROC Curve')
     plt.legend()
     # Plot Actual vs Predicted for Regression Task
     plt.subplot(1, 2, 2)
     sns.scatterplot(x=y_test_housing, y=y_pred_housing)
     plt.xlabel('Actual Values')
     plt.ylabel('Predicted Values')
     plt.title('Actual vs Predicted Housing Prices')
     plt.tight_layout()
     plt.show()
```



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[11]: y_pred_cancer=rf_classifier.predict(X_test_cancer)
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[9]: y_pred_housing=rf_regressor.predict(X_test_housing)

[10]: print("Predictions for Classification Task (Breast Cancer):",y_pred_cancer) print("Predictions for Classification Task (House Prediction):",y_pred_housing)