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# Import libraries
import pandas as pd
from sklearn.datasets import load_breast_cancer
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import classification_report, accuracy_score
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC
from sklearn.tree import DecisionTreeClassifier

# Load dataset
data = load_breast_cancer()
X = data.data
y = data.target

# Preprocessing - Scaling
scaler = StandardScaler()
X = scaler.fit_transform(X)

# Train-test split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

# Initialize models
models = {
    "K-Nearest Neighbors": KNeighborsClassifier(n_neighbors=5),
    "SVM": SVC(kernel='linear'),
    "Decision Tree": DecisionTreeClassifier(random_state=42)
}

# Train, predict, and evaluate
results = {}
for name, model in models.items():
    model.fit(X_train, y_train)
    y_pred = model.predict(X_test)

    accuracy = accuracy_score(y_test, y_pred)
    report = classification_report(y_test, y_pred, target_names=data.target_names, output_dict=True)

    results[name] = {
        "Accuracy": accuracy,
        "Precision": report['weighted avg']['precision'],
        "Recall": report['weighted avg']['recall'],
        "F1-score": report['weighted avg']['f1-score']
    }

# Display results
df_results = pd.DataFrame(results).T
print("\nPerformance Comparison:")
print(df_results)

# Detailed classification reports
for name, model in models.items():
    y_pred = model.predict(X_test)
    print(f"\n{name} Classification Report:\n")
    print(classification_report(y_test, y_pred, target_names=data.target_names))
```

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Performance Comparison:
                   Accuracy  Precision    Recall  F1-score
K-Nearest Neighbors  0.947368   0.947368  0.947368  0.947368
SVM                  0.956140   0.956488  0.956140  0.956237
Decision Tree        0.947368   0.947368  0.947368  0.947368

K-Nearest Neighbors Classification Report:

              precision    recall  f1-score   support

   malignant       0.93      0.93      0.93        43
      benign       0.96      0.96      0.96        71

    accuracy                           0.95       114
   macro avg       0.94      0.94      0.94       114
weighted avg       0.95      0.95      0.95       114


SVM Classification Report:

              precision    recall  f1-score   support

   malignant       0.93      0.95      0.94        43
      benign       0.97      0.96      0.96        71
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        accuracy                           0.96       114
       macro avg       0.95      0.96      0.95       114
    weighted avg       0.96      0.96      0.96       114


    Decision Tree Classification Report:

                  precision    recall  f1-score   support

       malignant       0.93      0.93      0.93        43
          benign       0.96      0.96      0.96        71

        accuracy                           0.95       114
       macro avg       0.94      0.94      0.94       114
    weighted avg       0.95      0.95      0.95       114
```