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In [1]: import pandas as pd
 import numpy as np
 from sklearn.datasets import load breast cancer
 from sklearn.model_selection import train_test_split
 from sklearn.preprocessing import StandardScaler
 from sklearn.linear_model import Perceptron
 from sklearn.metrics import accuracy_score, confusion_matrix, roc_curve, auc, precision_recall_curve
 import seaborn as sns
 import matplotlib.pyplot as plt
 data = load breast cancer()
 X = pd.DataFrame(data.data, columns=data.feature names)
 y = pd.Series(data.target)
 X_{\text{train}}, X_{\text{test}}, y_{\text{train}}, y_{\text{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)
 perceptron = Perceptron(max iter=1000, eta0=0.1, random state=42)
 perceptron.fit(X_train, y_train)
 y pred = perceptron.predict(X test)
 accuracy = accuracy_score(y_test, y_pred)
 print(f'Accuracy: {accuracy:.4f}')
 print(f'Final Weights: {perceptron.coef_}')
 conf_matrix = confusion_matrix(y_test, y_pred)
 plt.figure(figsize=(12, 4))
 plt.subplot(1, 3, 1)
 sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues',
              xticklabels=data.target_names,
              yticklabels=data.target_names)
 plt.xlabel('Predicted')
 plt.ylabel('True')
 plt.title('Confusion Matrix')
 fpr, tpr, _ = roc_curve(y_test, perceptron.decision_function(X_test))
 roc_auc = auc(fpr, tpr)
 plt.subplot(1, 3, 2)
 plt.plot(fpr, tpr, color='darkorange', lw=2, label=f'ROC curve (AUC = {roc_auc:.2f})')
 plt.plot([0, 1], [0, 1], color='navy', lw=2, linestyle='--')
 plt.xlabel('False Positive Rate')
 plt.ylabel('True Positive Rate')
 plt.title('ROC Curve')
 plt.legend(loc='lower right')
 precision, recall,
                       = precision recall curve(y test, perceptron.decision function(X test))
 plt.subplot(1, 3, 3)
 plt.plot(recall, precision, color='green', lw=2)
 plt.xlabel('Recall')
 plt.ylabel('Precision')
 plt.title('Precision-Recall Curve')
 plt.tight_layout()
 plt.show()
Accuracy: 0.9737
Final Weights: [[-0.29708989 0.3979993 -0.22971536 -0.36722303 -0.29458505 0.60765684
  -0.48742822 \ -0.87535615 \ \ 0.57180141 \ \ 0.38323832 \ -1.25108911 \ \ 0.09947304
  -0.20503338 - 1.0116695 - 1.12802674 0.30730166 0.40703831 - 0.82537704
   0.33157064 \quad 0.3003877 \quad -0.73262937 \quad -1.15607325 \quad -0.37557777 \quad -0.70380201
  -0.20128268 -0.03769604 -0.78013419 -0.92248386 -1.58936503 0.14415303]]
          Confusion Matrix
                                                         ROC Curve
                                                                                              Precision-Recall Curve
                                          1.0
                                                                                  1.00
                                   60
                                                                                  0.95
                                          0.8
                                   50
                                                                                  0.90
                                        Positive Rate
                                          0.6
                                                                                  0.85
                                    40
                                                                                  0.80
                                   30
                                          0.4
                                                                                  0.75
                                   20
                        69
                                                                                  0.70
                                          0.2
                                  - 10
                                                                                  0.65
                                                            ROC curve (AUC = 1.00)
```

0.0

0.0

0.4

False Positive Rate

0.6

0.8

1.0

0.0

0.2

0.4

Recall

0.6

0.8

1.0

malignant

Predicted

benign