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In [4]: import pandas as pd
        from sklearn.model selection import train test split
        from sklearn.ensemble import RandomForestClassifier
        from sklearn.metrics import accuracy score, confusion matrix, ConfusionMatrixD
        import matplotlib.pyplot as plt
        # --- Step 1: Load dataset from URL ---
        url = "https://raw.githubusercontent.com/jbrownlee/Datasets/master/pima-indiar
        # Column names as per dataset description
        columns = ["Pregnancies", "Glucose", "BloodPressure", "SkinThickness",
                   "Insulin", "BMI", "DiabetesPedigreeFunction", "Age", "Outcome"]
        df = pd.read csv(url, header=None, names=columns)
        print("Dataset preview:")
        print(df.head())
        # --- Step 2: Split into features and target ---
        X = df.drop("Outcome", axis=1)
        y = df["Outcome"]
        # --- Step 3: Train-test split ---
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25, rand
        # --- Step 4: Train Random Forest classifier ---
        rf = RandomForestClassifier(n_estimators=100, random state=42)
        rf.fit(X train, y train)
        # --- Step 5: Predictions & Evaluation ---
        y pred = rf.predict(X test)
        print("\nAccuracy:", accuracy_score(y_test, y_pred))
      Dataset preview:
         Pregnancies Glucose BloodPressure SkinThickness Insulin
                                                                     BMI \
      0
                   6
                          148
                                          72
                                                         35
                                                                  0 33.6
      1
                   1
                          85
                                          66
                                                         29
                                                                  0 26.6
                   8
                                          64
      2
                          183
                                                        0
                                                                 0 23.3
      3
                   1
                           89
                                          66
                                                         23
                                                                 94 28.1
      4
                   0
                          137
                                          40
                                                         35
                                                                168 43.1
         DiabetesPedigreeFunction Age Outcome
      0
                            0.627 50
                            0.351 31
      1
                                              0
      2
                            0.672 32
                                              1
      3
                            0.167
                                    21
                                              0
      4
                            2.288
                                    33
                                              1
      Accuracy: 0.734375
In [3]: # --- Step 6: Feature Importance ---
        importances = rf.feature importances
        features = X.columns
        indices = importances.argsort()[::-1]
```

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plt.figure(figsize=(10,6))
plt.title("Feature Importance")
plt.bar(range(X.shape[1]), importances[indices], align="center")
plt.xticks(range(X.shape[1]), [features[i] for i in indices], rotation=45)
plt.ylabel("Importance")
plt.tight_layout()
plt.show()
```

