Classifier Visualization

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You are currently looking at **version 1.0** of this notebook. To download notebooks and datafiles, as well as get help on Jupyter notebooks in the Coursera platform, visit the Jupyter Notebook FAQ course resource.

1 Classifier Visualization Playground

The purpose of this notebook is to let you visualize various classifiers' decision boundaries.

The data used in this notebook is based on the UCI Mushroom Data Set stored in mushrooms.csv.

In order to better vizualize the decision boundaries, we'll perform Principal Component Analysis (PCA) on the data to reduce the dimensionality to 2 dimensions. Dimensionality reduction will be covered in a later module of this course.

Play around with different models and parameters to see how they affect the classifier's decision boundary and accuracy!

```
In []: %matplotlib notebook
    import pandas as pd
    import numpy as np
    import matplotlib.pyplot as plt
    from sklearn.decomposition import PCA
    from sklearn.model_selection import train_test_split

    df = pd.read_csv('readonly/mushrooms.csv')
    df2 = pd.get_dummies(df)

    df3 = df2.sample(frac=0.08)

    X = df3.iloc[:,2:]
    y = df3.iloc[:,1]
pca = PCA(n_components=2).fit_transform(X)
```

```
X_train, X_test, y_train, y_test = train_test_split(pca, y, random_state=0)
        plt.figure(dpi=120)
        plt.scatter(pca[y.values==0,0], pca[y.values==0,1], alpha=0.5, label='Edib'
        plt.scatter(pca[y.values==1,0], pca[y.values==1,1], alpha=0.5, label='Poiso
        plt.legend()
        plt.title('Mushroom Data Set\nFirst Two Principal Components')
        plt.xlabel('PC1')
        plt.ylabel('PC2')
        plt.gca().set_aspect('equal')
In [ ]: def plot_mushroom_boundary(X, y, fitted_model):
            plt.figure(figsize=(9.8,5), dpi=100)
            for i, plot_type in enumerate(['Decision Boundary', 'Decision Probabil:
                plt.subplot (1, 2, i+1)
                mesh_step_size = 0.01 # step size in the mesh
                x_{min}, x_{max} = X[:, 0].min() - .1, <math>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, mesh_step_size), np.ar
                if i == 0:
                    Z = fitted model.predict(np.c [xx.ravel(), yy.ravel()])
                else:
                    try:
                        Z = fitted_model.predict_proba(np.c_[xx.ravel(), yy.ravel()
                    except:
                        plt.text(0.4, 0.5, 'Probabilities Unavailable', horizontal
                             verticalalignment='center', transform = plt.gca().transform
                        plt.axis('off')
                        break
                Z = Z.reshape(xx.shape)
                plt.scatter(X[y.values==0,0], X[y.values==0,1], alpha=0.4, label='H
                plt.scatter(X[y.values==1,0], X[y.values==1,1], alpha=0.4, label='H
                plt.imshow(Z, interpolation='nearest', cmap='RdYlBu_r', alpha=0.15,
                           extent=(x_min, x_max, y_min, y_max), origin='lower')
                plt.title(plot_type + '\n' +
                          str(fitted_model).split('(')[0]+ ' Test Accuracy: ' + str
                plt.gca().set_aspect('equal');
            plt.tight_layout()
            plt.subplots_adjust(top=0.9, bottom=0.08, wspace=0.02)
In [ ]: from sklearn.linear_model import LogisticRegression
        model = LogisticRegression()
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model.fit(X_train,y_train)
        plot_mushroom_boundary(X_test, y_test, model)
In [ ]: from sklearn.neighbors import KNeighborsClassifier
        model = KNeighborsClassifier(n_neighbors=20)
        model.fit(X_train,y_train)
        plot_mushroom_boundary(X_test, y_test, model)
In [ ]: from sklearn.tree import DecisionTreeClassifier
        model = DecisionTreeClassifier(max_depth=3)
        model.fit(X_train,y_train)
       plot_mushroom_boundary(X_test, y_test, model)
In [ ]: from sklearn.tree import DecisionTreeClassifier
        model = DecisionTreeClassifier()
        model.fit(X_train,y_train)
        plot_mushroom_boundary(X_test, y_test, model)
In [ ]: from sklearn.ensemble import RandomForestClassifier
        model = RandomForestClassifier()
        model.fit(X_train,y_train)
       plot_mushroom_boundary(X_test, y_test, model)
In [ ]: from sklearn.svm import SVC
        model = SVC(kernel='linear')
        model.fit(X_train,y_train)
        plot_mushroom_boundary(X_test, y_test, model)
In [ ]: from sklearn.svm import SVC
        model = SVC(kernel='rbf', C=1)
        model.fit(X_train,y_train)
        plot_mushroom_boundary(X_test, y_test, model)
In [ ]: from sklearn.svm import SVC
        model = SVC(kernel='rbf', C=10)
```

```
model.fit(X_train,y_train)

plot_mushroom_boundary(X_test, y_test, model)

In []: from sklearn.naive_bayes import GaussianNB

model = GaussianNB()
model.fit(X_train,y_train)

plot_mushroom_boundary(X_test, y_test, model)

In []: from sklearn.neural_network import MLPClassifier

model = MLPClassifier()
model.fit(X_train,y_train)

plot_mushroom_boundary(X_test, y_test, model)

In []:
```