PCA vs CCA

Here we look at an example comparing the classification of some data after projection using PCA and CCA techniques.

yy = a * xx - (clf.intercept_[0]) / w[1]
plt.plot(xx, yy, linestyle, label=label)

```
In [0]: import numpy as np
import matplotlib.pyplot as plt

from sklearn.datasets import make_multilabel_classification
from sklearn.multiclass import OneVsRestClassifier
from sklearn.svm import SVC
from sklearn.decomposition import PCA
from sklearn.cross_decomposition import CCA

In [0]: def plot_hyperplane(clf, min_x, max_x, linestyle, label):
    # get the separating hyperplane
    w = clf.coef_[0]
    a = -w[0] / w[1]
    xx = np.linspace(min_x - 5, max_x + 5) # make sure the line is long enough
```

1 of 3 1/12/20, 3:27 PM

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In [0]: def plot_subfigure(X, Y, subplot, title, transform):
            if transform == "pca":
               X = PCA(n_components=2).fit_transform(X)
            elif transform == "cca":
               X = CCA(n\_components=2).fit(X, Y).transform(X)
               raise ValueError
           min_x = np.min(X[:, 0])
           max_x = np.max(X[:, 0])
           min_y = np.min(X[:, 1])
           max_y = np.max(X[:, 1])
           classif = OneVsRestClassifier(SVC(kernel='linear'))
           classif.fit(X, Y)
           plt.subplot(2, 2, subplot)
           plt.title(title)
            zero_class = np.where(Y[:, 0])
            one_class = np.where(Y[:, 1])
           facecolors='none', linewidths=2, label='Class 1')
            plt.scatter(X[one_class, 0], X[one_class, 1], s=80, edgecolors='orange',
                       facecolors='none', linewidths=2, label='Class 2')
           plot_hyperplane(classif.estimators_[0], min_x, max_x, 'k--',
                           'Boundary\nfor class 1')
            plot_hyperplane(classif.estimators_[1], min_x, max_x, 'k-.',
                           'Boundary\nfor class 2')
            plt.xticks(())
           plt.yticks(())
            plt.xlim(min_x - .5 * max_x, max_x + .5 * max_x)
           plt.ylim(min_y - .5 * max_y, max_y + .5 * max_y)
            if subplot == 2:
               plt.xlabel('First principal component')
               plt.ylabel('Second principal component')
               plt.legend(loc="upper left")
```

2 of 3 1/12/20, 3:27 PM

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In [4]: plt.figure(figsize=(8, 6))

X, Y = make_multilabel_classification(n_classes=2, n_labels=1, allow_unlabeled=True, random_state=1)

plot_subfigure(X, Y, 1, "With unlabeled samples + CCA", "cca")
plot_subfigure(X, Y, 2, "With unlabeled samples + PCA", "pca")

X, Y = make_multilabel_classification(n_classes=2, n_labels=1, allow_unlabeled=False, random_state=1)

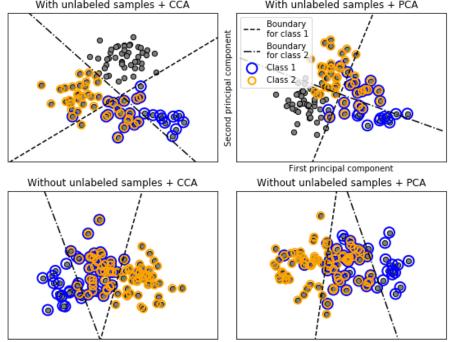
plot_subfigure(X, Y, 3, "Without unlabeled samples + CCA", "cca")
plot_subfigure(X, Y, 4, "Without unlabeled samples + PCA", "pca")

plt.subplots_adjust(.04, .02, .97, .94, .09, .2)
plt.show()

With unlabeled samples + CCA

With unlabeled samples + PCA

With unlabeled samples + PCA
```



Source: scikit-learn multilabel classification example

3 of 3 1/12/20, 3:27 PM