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# -*- coding: utf-8 -*-
```

```
"""
```

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Simple example of PCA (Principal Component Analysis) using the public Iris dataset.

```
"""
```

```
from sklearn.datasets import load_iris
from sklearn.decomposition import PCA
import pylab as pl
from itertools import cycle
```

```
iris = load_iris()
```

```
numSamples, numFeatures = iris.data.shape
```

```
#print(numSamples)
```

```
#print(numFeatures)
```

```
#print(list(iris.target_names))
```

```
#distilling the 4 dimensions down into 2
```

```
X = iris.data
```

```
pca = PCA(n_components=2, whiten=True).fit(X)
```

```
X_pca = pca.transform(X)
```

```
#print(pca.components_)
```

```
#print(pca.explained_variance_ratio_)
```

```
#print(sum(pca.explained_variance_ratio_))
```

```
#[0.92461872 0.05306648]
```

```
#0.977685206318795
```

```
#Although we have discarded two of our four dimensions, PCA has chosen the  
#remaining two dimensions so well that we've captured 92% of the variance in  
#our data in a single dimension alone. The second dimension just gives an  
#additional 5%; altogether by projecting it down to two dimensions, we've only  
#really lost less than 3% of the variance in the data
```

```
#Plotting the resultant points.
```

```
%matplotlib inline
```

```
from pylab import *
```

```
colors = cycle('rgb')
```

```
target_ids = range(len(iris.target_names))
```

```
pl.figure()
```

```
for i, c, label in zip(target_ids, colors, iris.target_names):
```

```
    pl.scatter(X_pca[iris.target == i, 0], X_pca[iris.target == i, 1],  
              c=c, label=label)
```

```
pl.legend()
```

```
pl.show()
```

#The three different types of Iris are still clustered fairly well.

