# **Principal Component Analysis**

#### Goal

PCA finds a new set of dimensions such that all the dimensions are orthogonal (and hence linearly independent) and ranked according to the variance of data along them.

Find a transformation such that

- · The tranformed features are linearly independent
- Dimensionality can be reduced by taking only the dimensions with the highest importance
- · Those newly found dimensions should minimize the projection error
- The projected points should have maximum spread, i.e. maximum variance

### **Variance**

How much variation or spread the data has.

$$Var(X) = \frac{1}{n} \sum_{i} (X_i - \bar{X})^2$$

#### **Covariance Matrix**

Indicates the level to which two variables vary together.

$$Cov(X,Y) = \frac{1}{n} \sum_{i} (X_i - \bar{X})(Y_i - \bar{Y})^T$$

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## Eigenvector, Eigenvalues

The eigenvectors point in the direction of the maximum variance, and the corresponding eigenvalues indicates the importance of its corresponding eigen vector.

$$\overrightarrow{Av} = \lambda \overrightarrow{v}$$

#### Approach

- -Subtract the mean from X
- -Calculate Cov(X,X)
- -Calculate eigenvectors and eigenvalues of covariance matrix
- -Sort the eigenvectors according to their eigenvalues in decreasing order
- -Choose first k eigenvectors and that will be the new k dimensions
- -Transform the original n dimensional data points into k dimensions (=Projections with dot product)

#### Using Iris test data from SKlearn

```
#PRINCIPAL COMPONENT ANALYSIS IN PYTHON
import numpy as np

##---- Begin PCA Implementation
class PCA:
    def __init__(self,n_components):
        self.n_components = n_components
        self.components = None
        self.mean = None

    def fit(self,X): #X is the data to be transformed
    #sort eigenvectors
    #store first n eigenvectors

#mean
    self.mean = np.mean(X,axis=0)
```

```
X = X - self.mean
        #covariance
        #row = 1 sample, columns=features so we T transpose
        cov = np.cov(X.T) #feature
        eigenvalues,eigenvectors = np.linalg.eig(cov) #return as col vectors
        #v[:,i]
        #sort eigenvectors
        eigenvectors = eigenvectors.T #transformed
        #sort in descendig order using -1
        idxs = np.argsort(eigenvalues)[::-1]
        eigenvalues = eigenvalues[idxs]
        eigenvectors = eigenvectors[idxs]
        #store first n eigenvectors
        self.components = eigenvectors[0:self.n_components]
    def transform(self,X):
        #project data
       X = X - self.mean
        return np.dot(X,self.components.T)
##---- End PCA Implementation
from sklearn import datasets
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
data = datasets.load iris()
```

```
X = data.data
y = data.target
#we have 105 samples and four features
pca = PCA(2)
pca.fit(X)
X_projected = pca.transform(X)
print('Shape of X: ', X.shape)
print('Shape of transformed X: ', X_projected.shape)
x1 = X_projected[:,0]
x2 = X_projected[:,1]
plt.scatter(x1,x2,c=y, edgecolors='none',alpha=0.8,cmap=plt.cm.get_cmap('viridis',3))
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.colorbar()
plt.show()
```





