Dimensionality reduction techniques: Part 1

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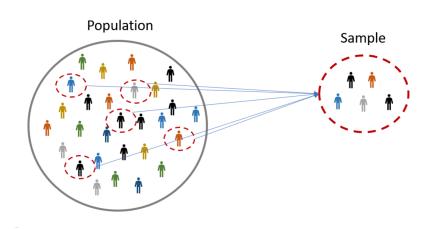
Motivation

Why dimensionality reduction?

Important "toolkit" in machine learning:

- Preprocessing step for other algorithms (reduce size)
- Visualization
- Compression
- Interpolation

Population vs sample



Population vs sample

Population mean

$$\mu = \frac{\sum_{i=1}^{N} x_i}{N} \tag{1}$$

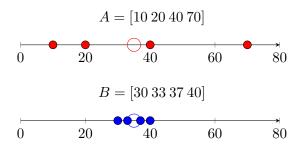
N is number of items in the population

Sample mean

$$\bar{X} = \frac{\sum_{i=1}^{n} x_i}{n} \tag{2}$$

n is number of items in the sample

Let's take a look on two samples:



Here, $\bar{A}=\bar{B}=35.$ Unfortunately, mean doesn't tell us a lot except for a middle point.

For our two sets, $A = [10\ 20\ 40\ 70]$ and $B = [30\ 33\ 37\ 40]$, we would be more interested in the *spread* of the data. So, how do we calculate it?

Standard deviation

$$s = \sqrt{\frac{\sum_{i=1}^{n} (X_i - \bar{X})^2}{(n-1)}}$$
 (3)

In plain English, it is the "average distance from the mean of the data set to a point."

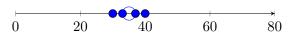
Set 1:
$$A = [10\ 20\ 40\ 70]$$
, and $\bar{A} = 35$



Let's calculate standard deviation:

\overline{A}	$(A - \bar{A})$	$(A-\bar{A})^2$
10	-25	625
20	-15	225
40	5	25
70	35	1,225
Total		2,100
Divided by (n-1)		700
Square root		26.4575

Set 2:
$$B = [30 \ 33 \ 37 \ 40]$$
, and $\bar{B} = 35$



Let's calculate standard deviation:

\overline{B}	$(B-\bar{B})$	$(B-\bar{B})^2$
30	-5	25
33	-2	4
37	2	4
40	5	25
Total		58
Divided by (n-1)		19.333
Square root		4.397

Variance

Similar to standard deviation So, how do we calculate it?

Variance

$$s^{2} = \frac{\sum_{i=1}^{n} (X_{i} - \bar{X})^{2}}{(n-1)}$$
 (4)

Almost identical to the standard deviation (there is no square root in the formula).

Covariance

Previous measures are 1-dimensional. How do we check relationship between the dimensions?

Covariance

Variance

$$var(X) = \frac{\sum_{i=1}^{n} (X_i - \bar{X})(X_i - \bar{X})}{(n-1)}$$
 (5)

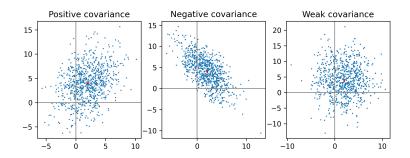
Covariance

$$covar(X,Y) = \frac{\sum_{i=1}^{n} (X_i - \bar{X})(Y_i - \bar{Y})}{(n-1)}$$
 (6)

How much dimensions vary from the mean with respect to each other.

Covariance between one dimension and itself gives variance.

Covariance



Covariance

Exercise: Find covariance between 2-dimensional dataset

Item number:	1	2	3	4	5
X	10	39	19	23	28
у	43	13	32	21	20

Ans: -120.55

Covariance matrix

When we have more than two dimensions, covariances between individual dimensions could be described in covariance matrix

Covariance matrix

$$C^{n \times n} = (c_{i,j} : c_{i,j} = cov(Dim_i, Dim_j))$$
(7)

where $C^{m\times n}$ is a matrix with n rows and m columns. Example for three dimensions:

$$c = \begin{pmatrix} cov(x, x) & cov(x, y) & cov(x, z) \\ cov(y, x) & cov(y, y) & cov(y, z) \\ cov(z, x) & cov(z, y) & cov(z, z) \end{pmatrix}$$
(8)

Since cov(a,b) = cov(b,a), the matrix is symmetrical about the main diagonal.

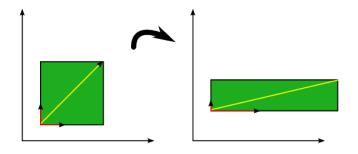
Covariance matrix

Exercise: Find covariance matrix for the 3-dimensional dataset

Item number:	1	2	3
X	1	-1	4
у	2	1	3
Z	1	3	-1

Ans:
$$\begin{pmatrix} 4.222 & 1.667 & -3.333 \\ 1.667 & 0.667 & -1.333 \\ -3.333 & -1.333 & 2.667 \end{pmatrix}$$

Eigenvectors



• Eigenvectors (red) do not change direction when a linear transformation (e.g. scaling) is applied to them. Other vectors (yellow) do.

Eigenvectors

An example of non-eigenvector:

$$\begin{pmatrix} 2 & 3 \\ 2 & 1 \end{pmatrix} \times \begin{pmatrix} 1 \\ 3 \end{pmatrix} = \begin{pmatrix} 11 \\ 5 \end{pmatrix} \tag{9}$$

An example of eigenvector:

$$\begin{pmatrix} 2 & 3 \\ 2 & 1 \end{pmatrix} \times \begin{pmatrix} 3 \\ 2 \end{pmatrix} = \begin{pmatrix} 12 \\ 8 \end{pmatrix} = 4 \times \begin{pmatrix} 3 \\ 2 \end{pmatrix} \tag{10}$$

Eigenvalues

The amount by which the original vector was scaled after multiplication by the square matrix

$$\begin{pmatrix} 2 & 3 \\ 2 & 1 \end{pmatrix} \times \begin{pmatrix} 3 \\ 2 \end{pmatrix} = \begin{pmatrix} 12 \\ 8 \end{pmatrix} = \boxed{4} \times \begin{pmatrix} 3 \\ 2 \end{pmatrix} \tag{11}$$

PCA

What is Principal Component Analysis?

A way of identifying patterns in data: by highlighting their similarities and differences [2]:

Pros

• Once patterns in the data, and you compress the data, ie. by reducing the number of dimensions, without much loss of information.

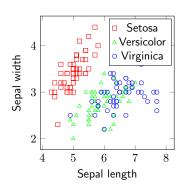
Cons

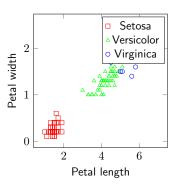
• Linear : difficult to unfold single ambiguous object really belongs in several disparate locations in the low-dimensional space.

Iris flower dataset

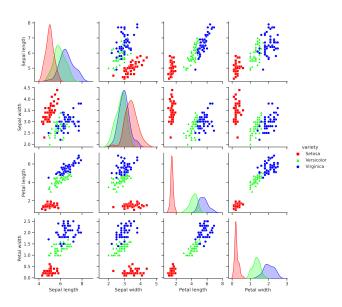


Iris flower dataset





Iris flower dataset



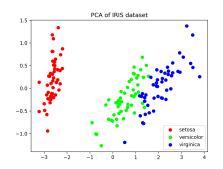
Applying PCA to Iris flower dataset

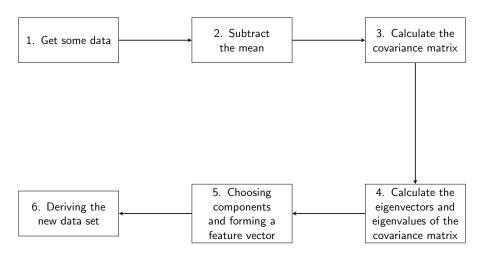
PCA automatically finds principal components (eg. *axes*) in feature space. Here, we defined two, so we can visualize in 2D easily.

```
from sklearn import datasets
from sklearn.decomposition import PCA
iris = datasets.load_iris()

X = iris.data
y = iris.target
target_names = iris.target_names

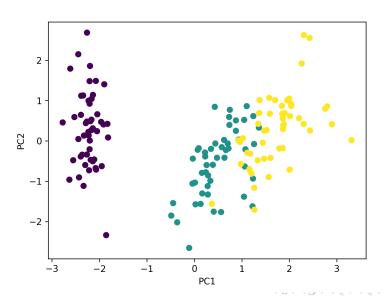
pca = PCA(n_components=2)
X_r = pca.fit(X).transform(X)
```





```
import numpy as np
from numpy.linalg import eig
def PCA_numpy(X,y):
   def mean(x): # np.mean(X, axis = 0)
        return sum(x)/len(x)
   def std(x): # np.std(X, axis = 0)
        return (sum((i - mean(x))**2 for i in x)/len(x))**0.5
   def Standardize data(X):
        return (X - mean(X))/std(X)
   def covariance(x):
        return (x.T @ x)/(x.shape[0]-1)
    # Step 1: Standardize the data
   X std = Standardize data(X)
    # Step 2: Find the covariance matrix
    cov_mat = covariance(X_std) # np.cov(X_std.T)
    # Step 3: Find the eigenvectors and eigenvalues of the covariance matrix
   eig_vals, eig_vecs = eig(cov_mat)
```

```
def continued ...
   max_abs_idx = np.argmax(np.abs(eig_vecs), axis=0)
    signs = np.sign(eig_vecs[max_abs_idx, range(eig_vecs.shape[0])])
   eig_vecs = eig_vecs*signs[np.newaxis,:]
   eig_vecs = eig_vecs.T
    # Step 4: Rearrange the eigenvectors and eigenvalues
   eig_pairs = [(np.abs(eig_vals[i]), eig_vecs[i,:]) for i in range(len(eig_vals))
    # Then, we sort the tuples from the highest to the lowest based on eigenvalues m
   eig_pairs.sort(key=lambda x: x[0], reverse=True)
   eig_vals_sorted = np.array([x[0] for x in eig_pairs])
   eig_vecs_sorted = np.array([x[1] for x in eig_pairs])
    # Step 5: Choose principal components
   k = 2
   W = eig_vecs_sorted[:k, :] # Projection matrix
    # Step 6: Project the data
   X_proj = X_std.dot(W.T)
   return X_proj
```



t-SNE (Stochastic neighbor embedding)

To be continued next time, Dimensionality reduction techniques: Part 2

Thank you for your attention!

Slides:

https://github.com/CodeSeoul/machine-learning/blob/master/ 221210-pca/lecture.pdf

- Recorded video: https://www.youtube.com/watch?v=UfwQyWDOKRM
- Solutions for exercises:
 https://github.com/CodeSeoul/machine-learning/blob/master/
 221210-pca/excel_ex.xlsx
- PCA from scratch: https://github.com/CodeSeoul/machine-learning/blob/master/ 221210-pca/practicum.ipynb
- Follow-up QA?
 http://discord.com/users/tuttelikz

References



Howard Anton and Chris Rorres. *Elementary linear algebra: applications version*. John Wiley & Sons, 2013.



Lindsay I Smith. "A tutorial on principal components analysis". In: (2002).