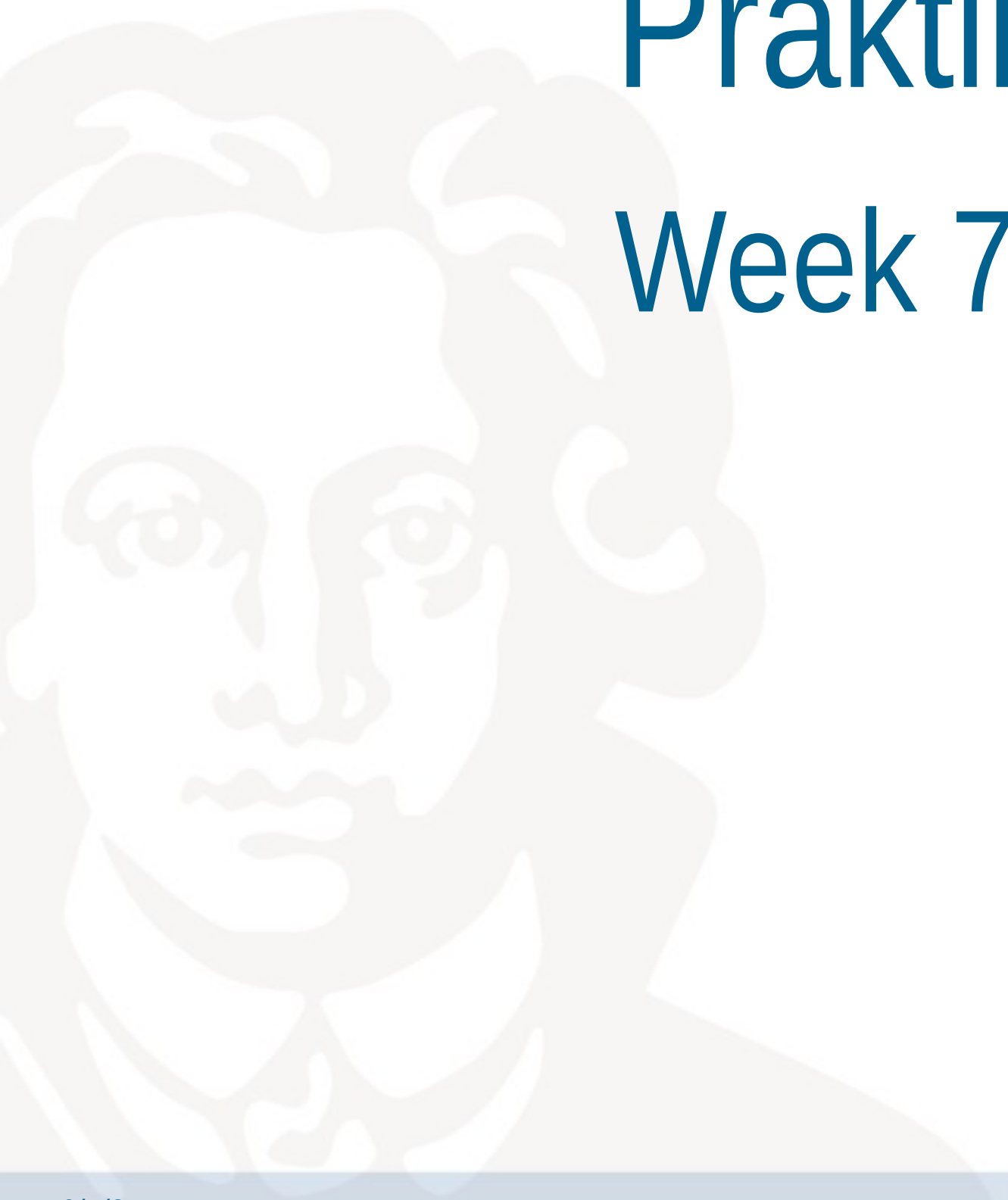


Martin Mundt, Dr. Iuliia Pliushch, Prof. Dr. Visvanathan Ramesh

# Pattern Analysis & Machine Intelligence

## Praktikum: MLPR-SS21

Week 7: K-Means, PCA and ICA

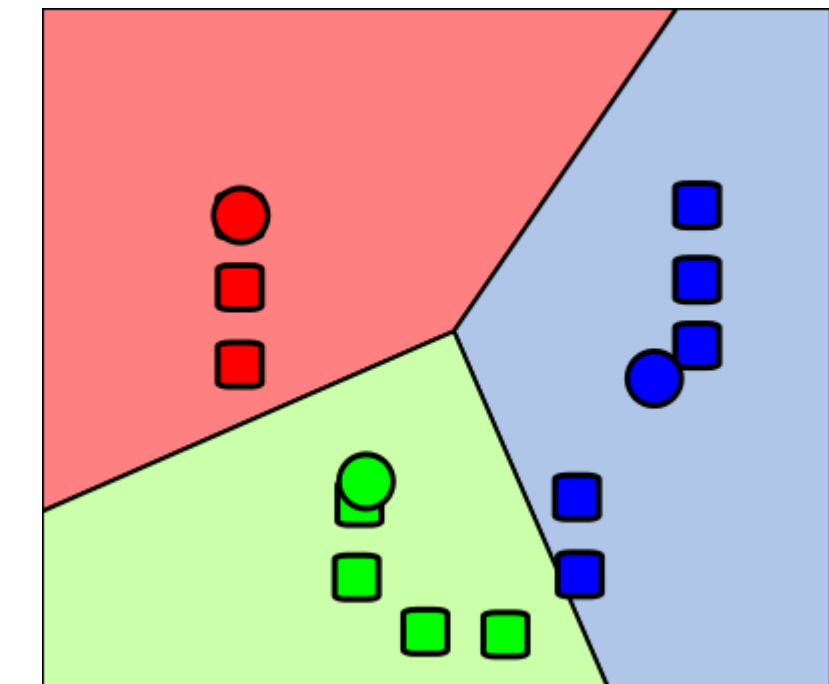
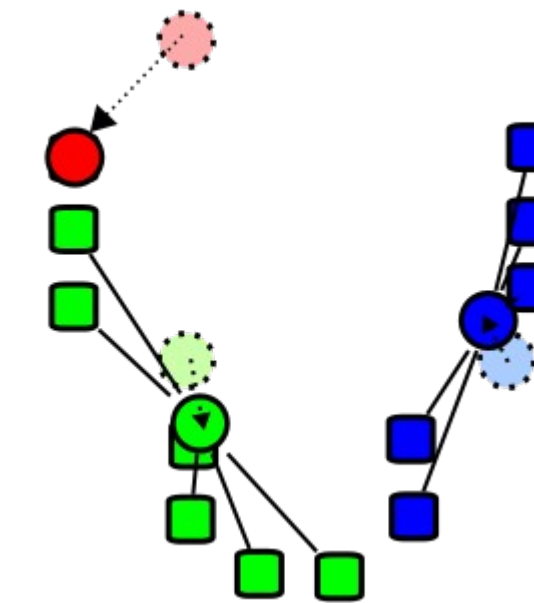
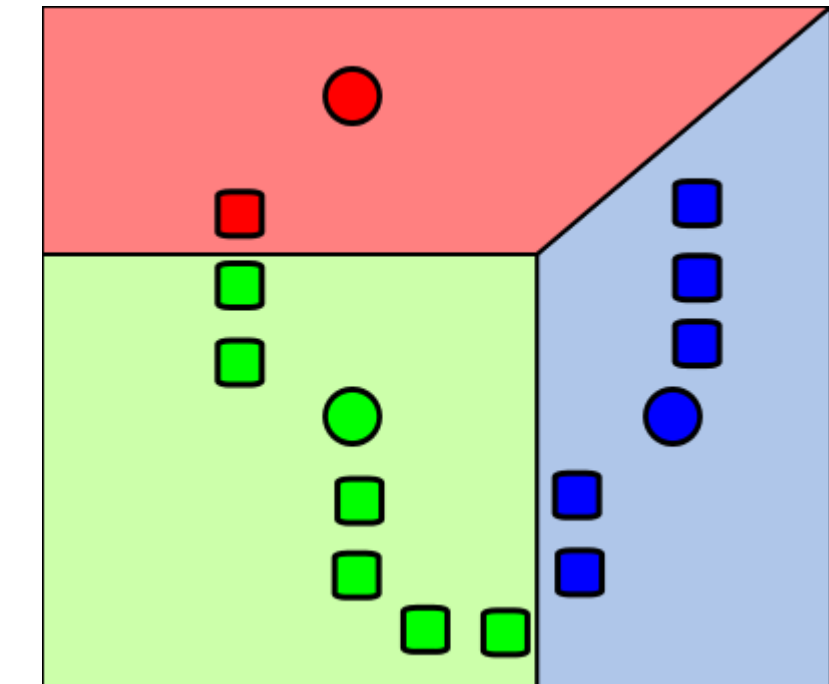
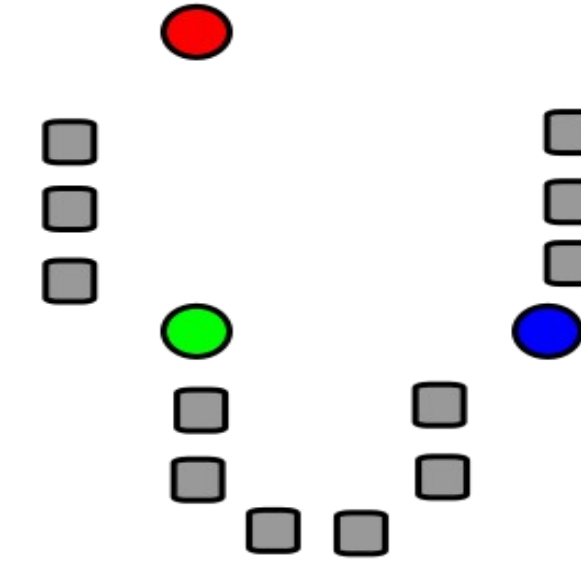


# Unsupervised learning: K-Means, PCA and ICA

- **K-means**: clustering algorithm which operates on the distances between points and the supposed cluster centers
- **PCA**: helps in analyzing the data variance and can be used for **data compression**
- **ICA**: separating a multivariate signal into non-Gaussian, **independent** subcomponents

# K-Means clustering (Lloyd algorithm)

- Input: d-dimensional data points
- Randomly initialize k cluster means
- Assign points to its closest cluster mean
- Update the cluster means and repeat the two previous steps until the means converge



<https://de.coursera.org/lecture/genomic-data/the-lloyd-algorithm-for-k-means-clustering-3O9eh>

<https://de.wikipedia.org/wiki/K-Means-Algorithmus>

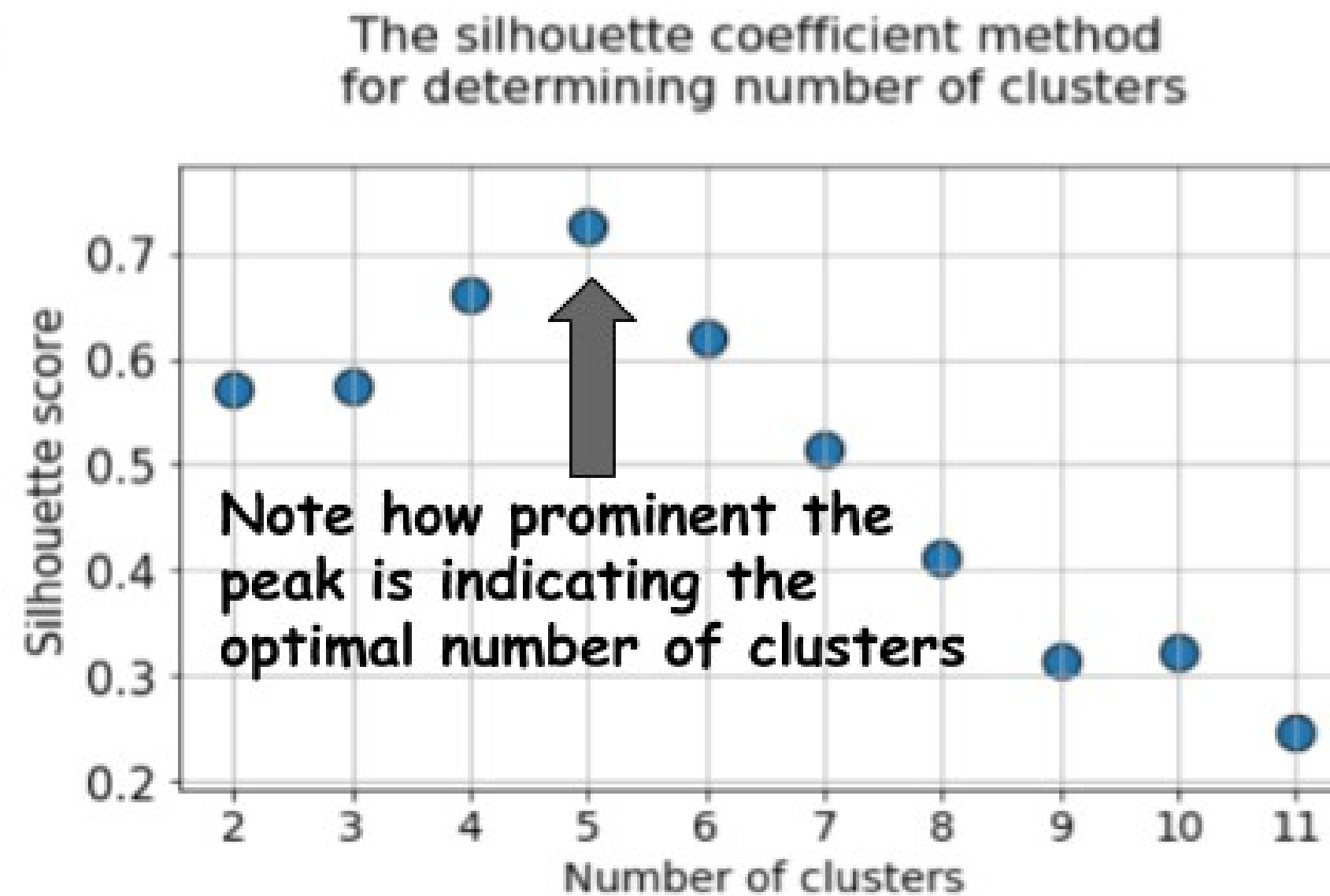
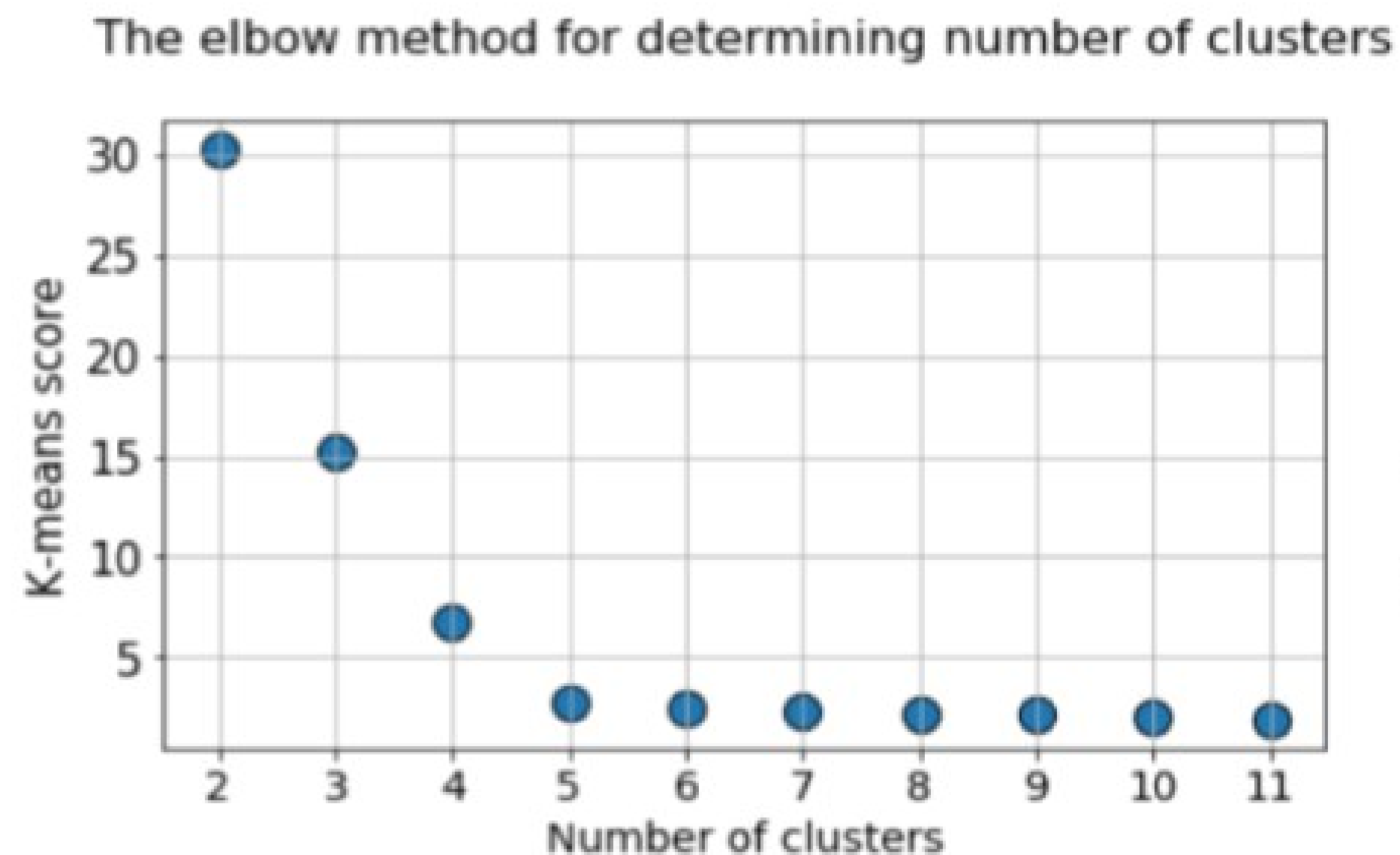
# K-Means: Difficulties

- How do we determine **k** – the number of clusters to split the data into?

Elbow	Silhouette
<ul style="list-style-type: none"> <li>- run k-means for different values of k</li> <li>- calculate <b>WCSS</b>: within cluster sum of squares</li> <li>- plot WCSS for growing k</li> <li>- take the k where 'the elbow bends'</li> </ul>	<ul style="list-style-type: none"> <li>- run k-means for different values of k</li> <li>- calculate the <b>average silhouette</b></li> <li>- plot the measure for growing k</li> <li>- take the k at the peak</li> </ul>

# K-Means: Difficulties (1)

- How do we determine  $k$  – the number of clusters to split the data into?

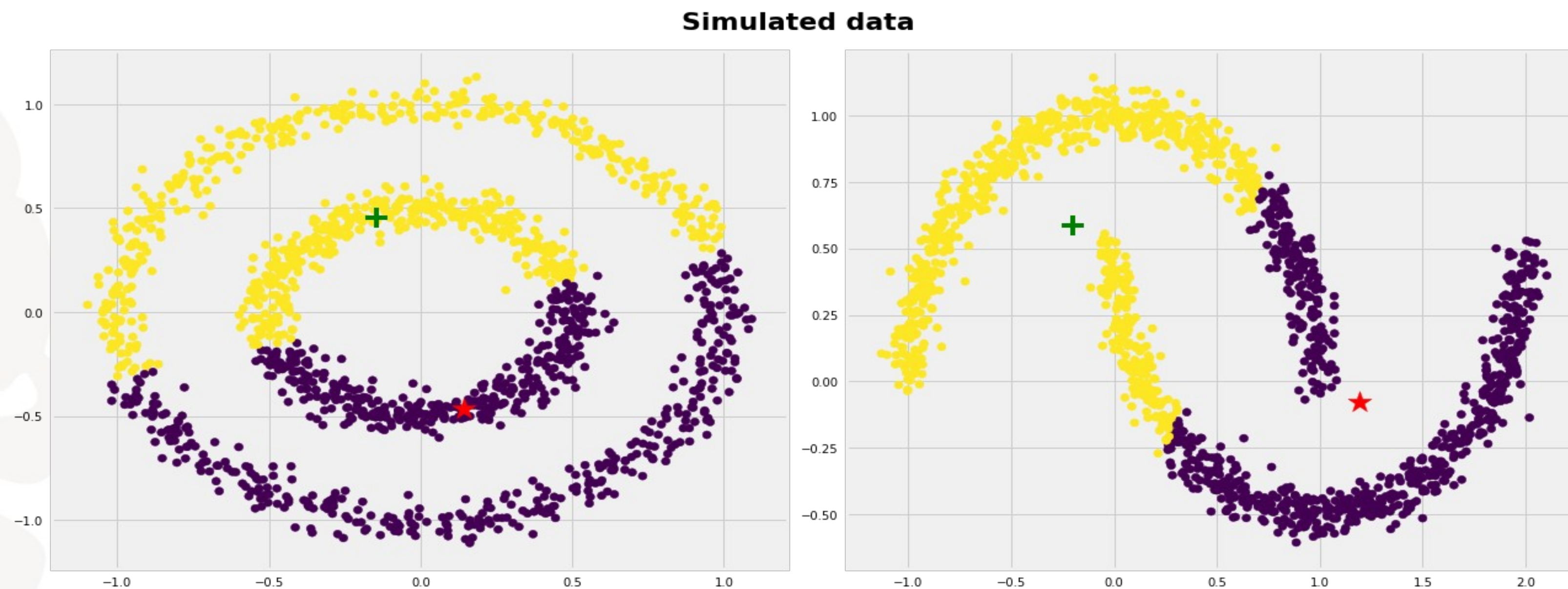


<https://towardsdatascience.com/clustering-metrics-better-than-the-elbow-method-6926e1f723a6>



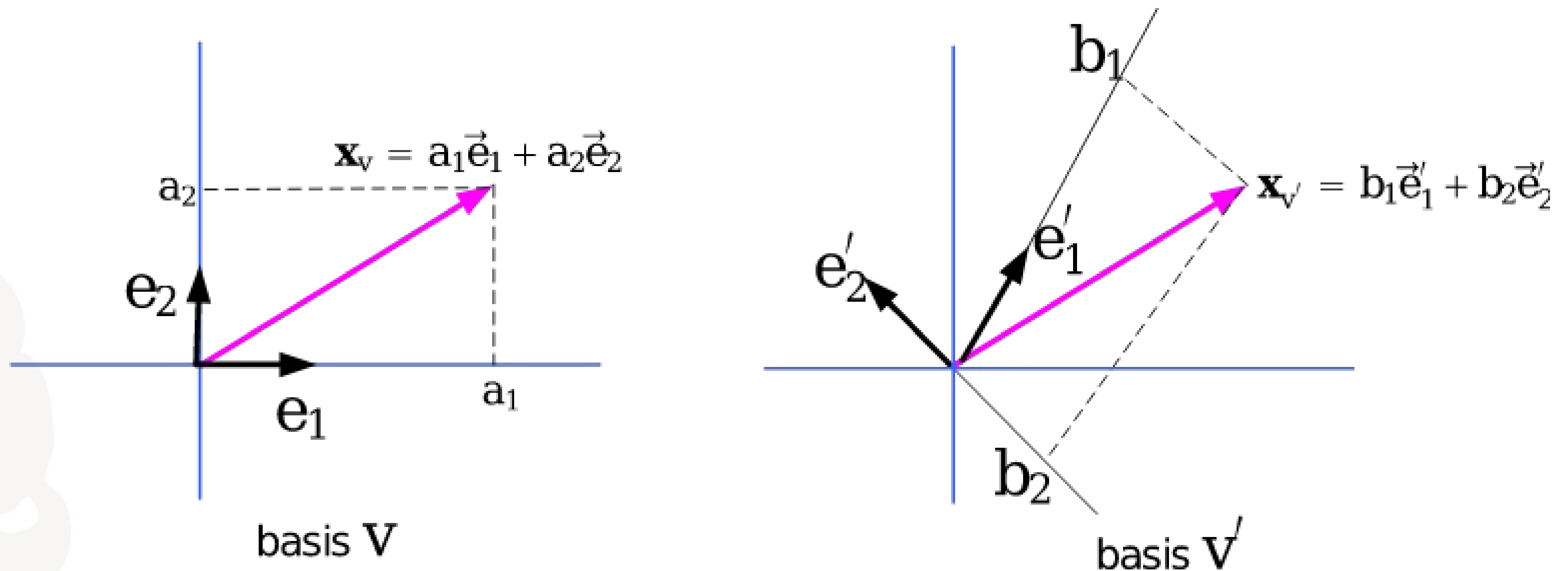
# K-Means: Difficulties (2)

- What if the cluster are not of a spherical shape?



<https://towardsdatascience.com/k-means-clustering-algorithm-applications-evaluation-methods-and-drawbacks-aa03e644b48a>

# PCA: Basis transformation



The same vector having different representation depending on basis used

[https://www.12000.org/my\\_notes/similarity\\_transformation\\_and\\_SVD/index.htm](https://www.12000.org/my_notes/similarity_transformation_and_SVD/index.htm)

# PCA: Eigenvalues & eigenvectors

eigenvalue

eigenvector

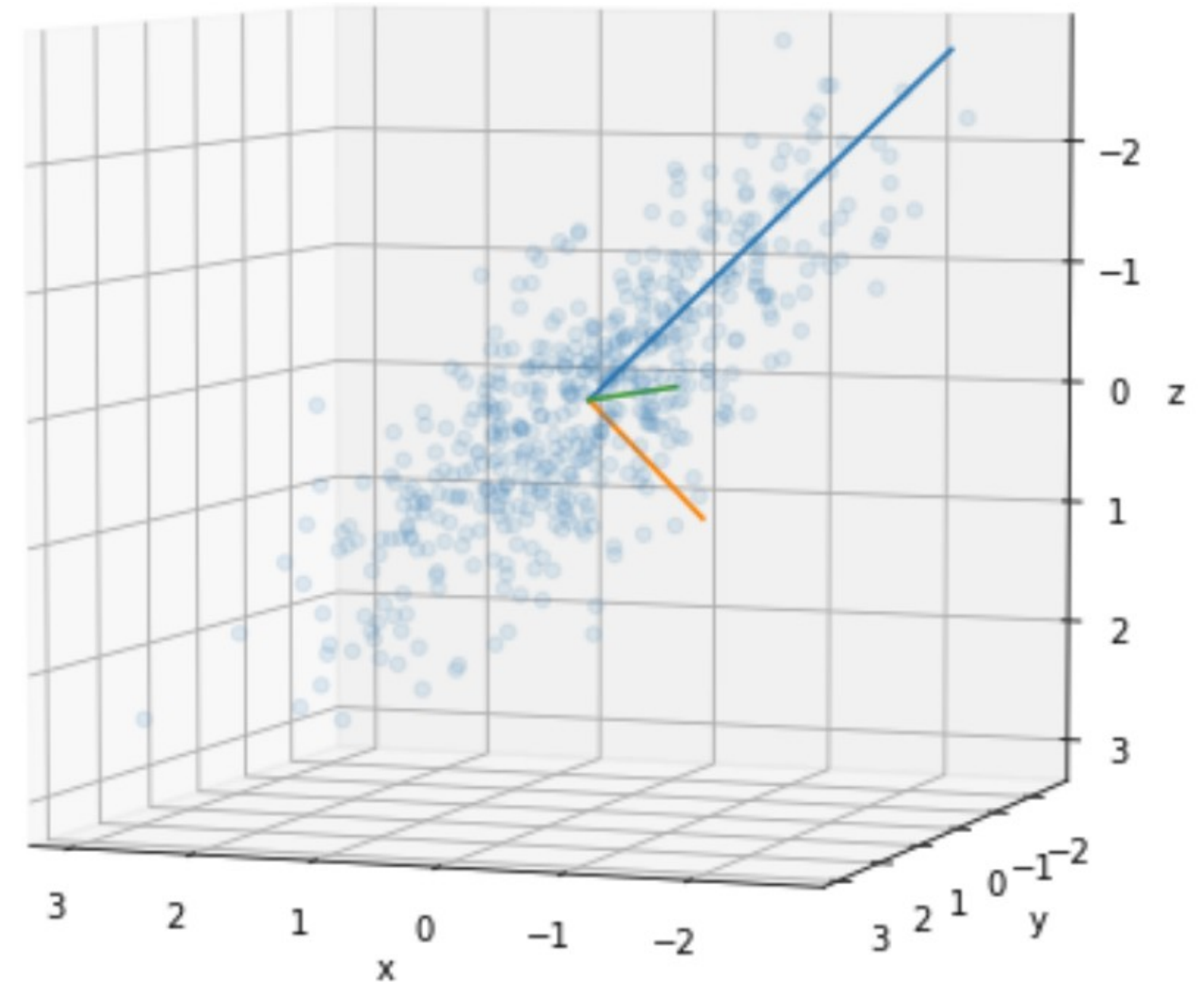
$$Av = \lambda v$$

Interpretation: the eigenvector  $v$  does not change (its direction) when multiplied by  $A$ , it is only scaled.



# PCA – Principal Component Analysis

- Input: d-dimensional data
- Subtract the mean from your data
- Compute the covariance matrix for your zero-mean data
- Compute the eigenvalues and eigenvectors of the **covariance matrix**
- Sort the **eigenvectors** (=principal components) in descending order according to the eigenvalues
- Pick a subset of them and transform your data



[http://www.iro.umontreal.ca/~pift6080/H09/documents/papers/pca\\_tutorial.pdf](http://www.iro.umontreal.ca/~pift6080/H09/documents/papers/pca_tutorial.pdf)

# Covariance matrix

- **Variance:**

$$\text{var}(X) = \frac{\sum_{i \in N} (x_i - \mu_x)^2}{N - 1} = \frac{\sum_{i \in N} (x_i - \mu_x) * (x_i - \mu_x)}{N - 1}$$

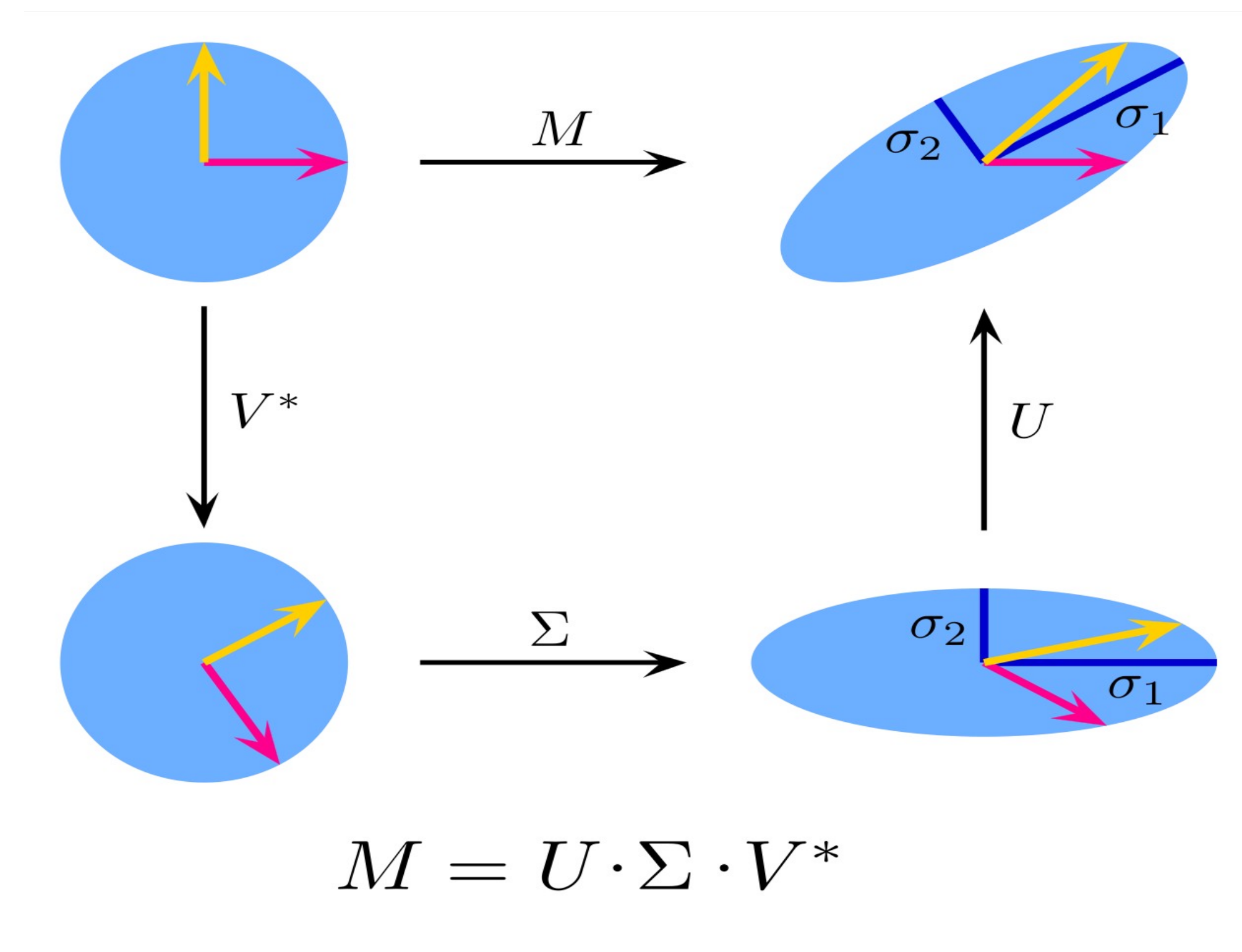
- **Covariance:**

$$\text{covar}(X, Y) = \frac{\sum_{i \in N} (x_i - \mu_x) * (y_i - \mu_y)}{N - 1}$$

[http://www.iro.umontreal.ca/~pift6080/H09/documents/papers/pca\\_tutorial.pdf](http://www.iro.umontreal.ca/~pift6080/H09/documents/papers/pca_tutorial.pdf)

# PCA: Compute the eigenvalues, -vectors

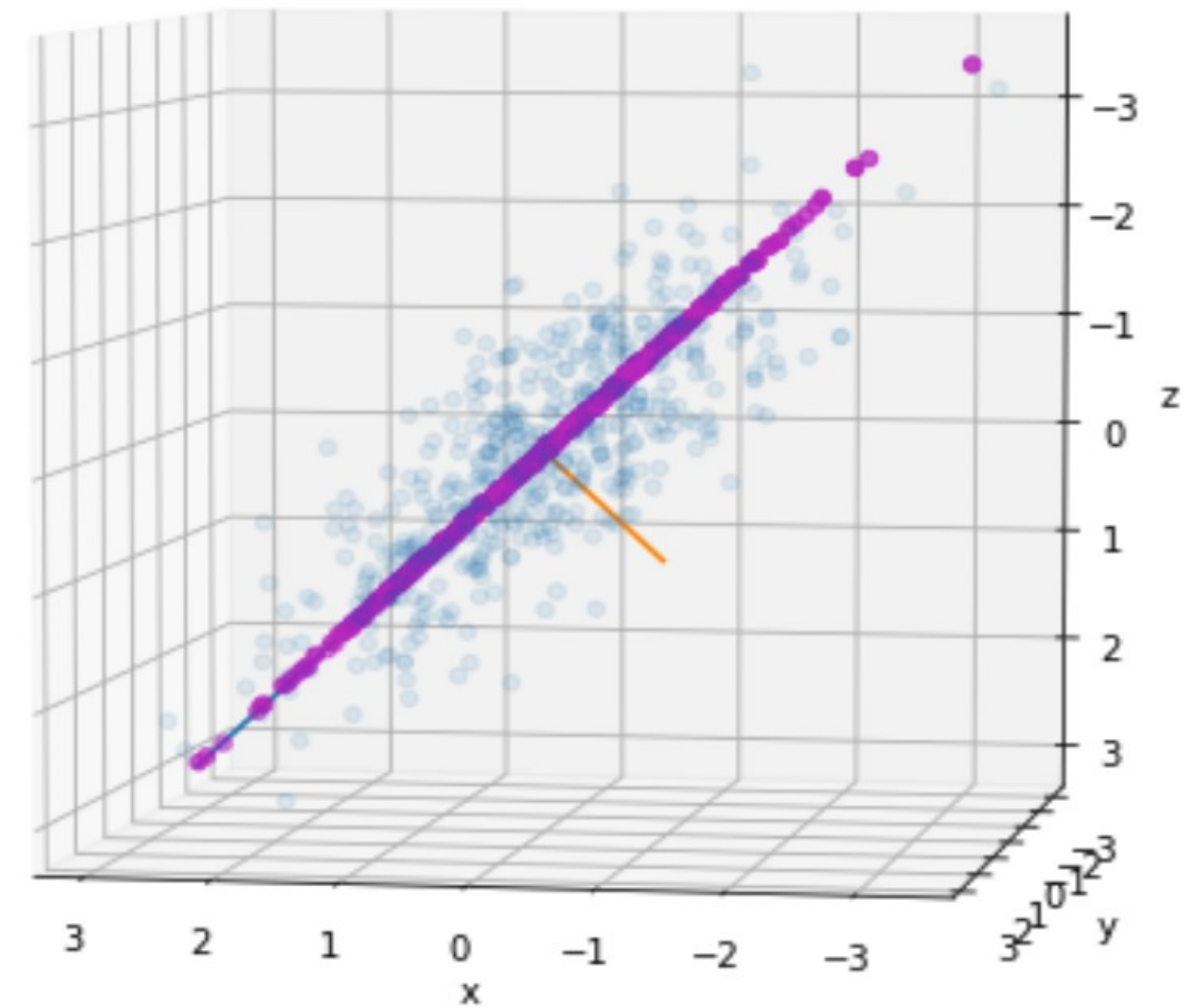
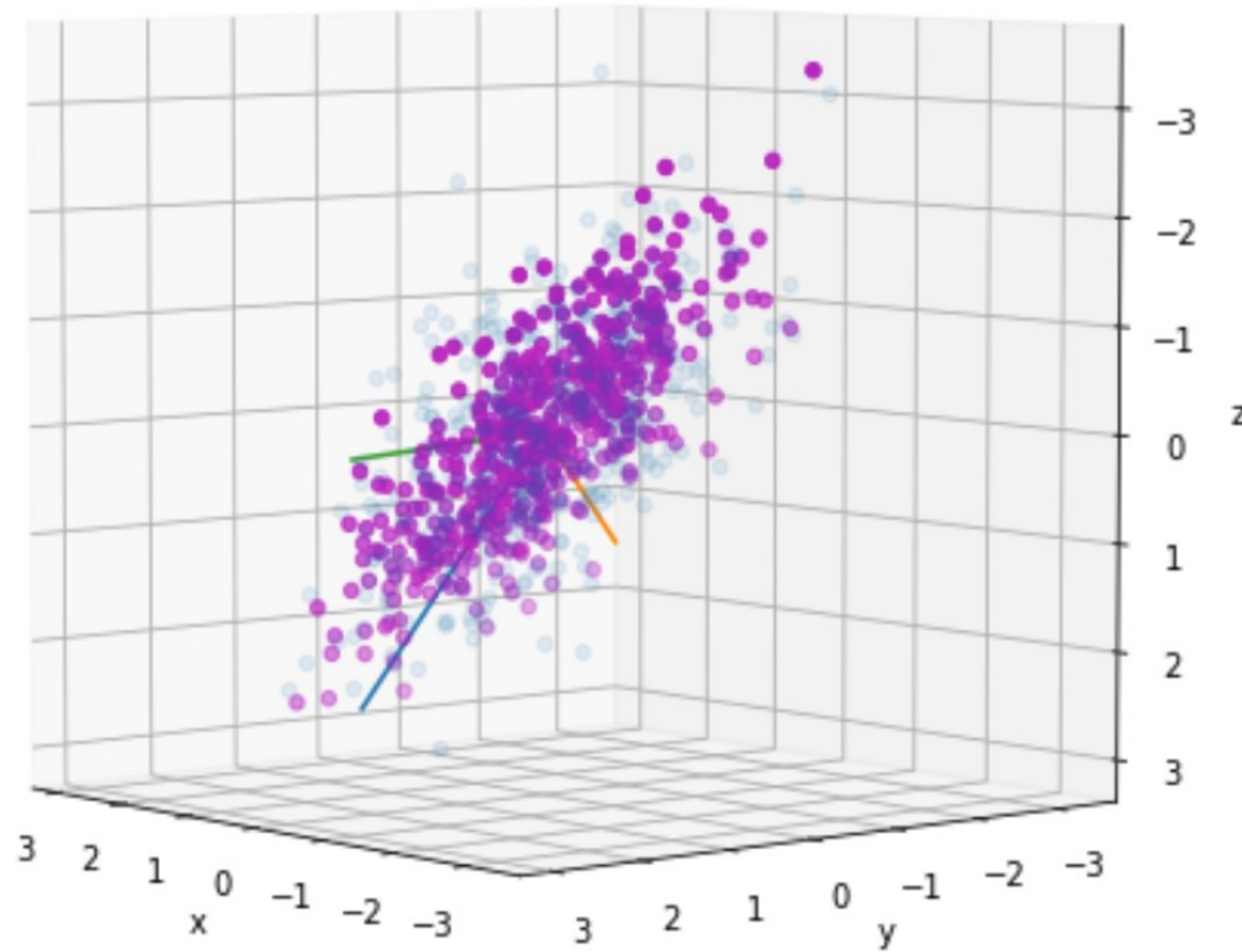
**SVD (singular value decomposition)** as a generalization of **eigendecomposition** (which is only applicable to square matrices)



[https://en.wikipedia.org/wiki/Singular\\_value\\_decomposition](https://en.wikipedia.org/wiki/Singular_value_decomposition)

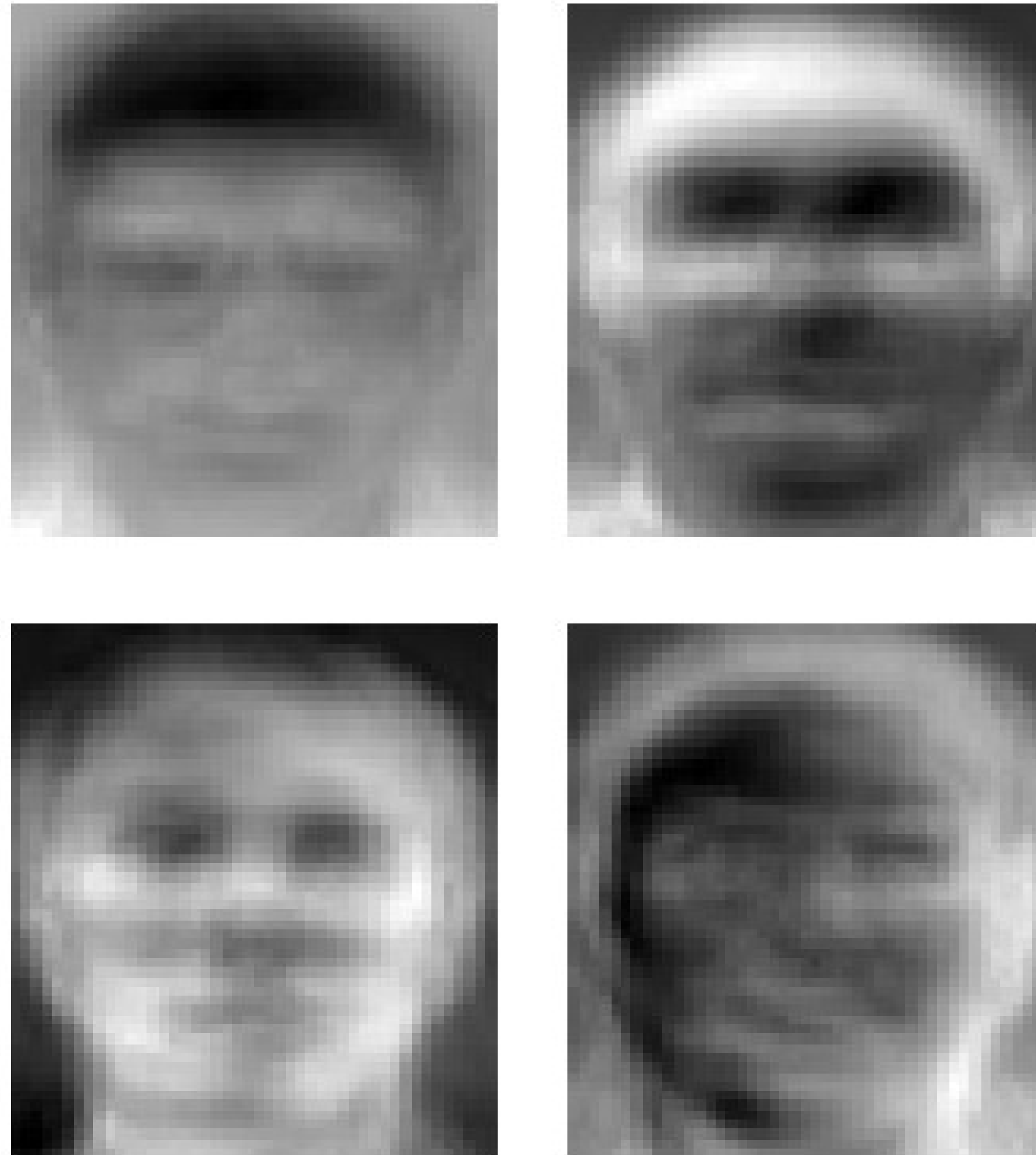


# Dimensionality reduction: 3D - 2D





# Facial recognition: Eigenfaces



<https://en.wikipedia.org/wiki/Eigenface>

# ICA – Independent component analysis

- separate a signal into **non-Gaussian** and **statistically independent** subcomponents



<https://edtech.engineering.utoronto.ca/files/item-76-cocktail-party-effectpng>

# ICA – Independent component analysis

1. Center  $\mathbf{x}$  by subtracting the mean
2. Whiten  $\mathbf{x}$  (PCA or ZCA)

## -----**fastICA**-----

3. Choose a random initial value for the de-mixing matrix  $\mathbf{W}$
4. Calculate the **new value** for  $w$
5. **Normalize**  $w$
6. Check whether algorithm has converged and if it hasn't, return to step 4

- ## -----
7. Take the **dot product of  $\mathbf{W}$  and  $\mathbf{x}$**  to get the independent source signals

<https://towardsdatascience.com/independent-component-analysis-ica-in-python-a0ef0db0955e>



# ICA: Whitening (2)

- **Reminder:** covariance matrix (**C**) of a symmetric positive semi-definite matrix can be decomposed into a diagonal matrix **D** and a transformation matrix **E**

$$C = E * D * E^T$$

- The transformation matrix for **PCA-whitening** is  $W_{PCA} = D^{1/2} * E^T$

- For **ZCA-whitening** is  $W_{ZCA} = E * D^{1/2} * E^T$

<https://stats.stackexchange.com/questions/117427/what-is-the-difference-between-zca-whitening-and-pca-whitening>



# ICA: fastICA (3-6)

- we search for such a matrix that the **independent sources**  $s = Wx$
- take as  $w$  a vector that maximizes the non-Gaussianity of  $w^T x$
- several measures of non-Gaussianity, for example **negentropy** using higher-order moments
- **decorrelate** outputs  $w_1^T x, \dots, w_n^T x$  so that no 2 vectors converge to the same maxima
- **convergence** when old and new  $w$  point into the same direction (their dot-product is 1)

A.Hyvärinen, E. Oja (2000). Independent component analysis: algorithms and applications. Neural Networks 13, 411-430

[https://itb.biologie.hu-berlin.de/~kempter/Teaching/2006\\_SS/hyvarinen\\_oja\\_2000.pdf](https://itb.biologie.hu-berlin.de/~kempter/Teaching/2006_SS/hyvarinen_oja_2000.pdf)

# ICA application: Image superposition



[https://upload.wikimedia.org/wikipedia/commons/7/74/Major\\_Mitchell%27s\\_Cockatoo\\_1\\_-\\_Mt\\_Grenfell.jpg](https://upload.wikimedia.org/wikipedia/commons/7/74/Major_Mitchell%27s_Cockatoo_1_-_Mt_Grenfell.jpg)

[https://upload.wikimedia.org/wikipedia/commons/7/73/Lion\\_waiting\\_in\\_Namibia.jpg](https://upload.wikimedia.org/wikipedia/commons/7/73/Lion_waiting_in_Namibia.jpg)

# Difference between PCA and ICA

## PCA:

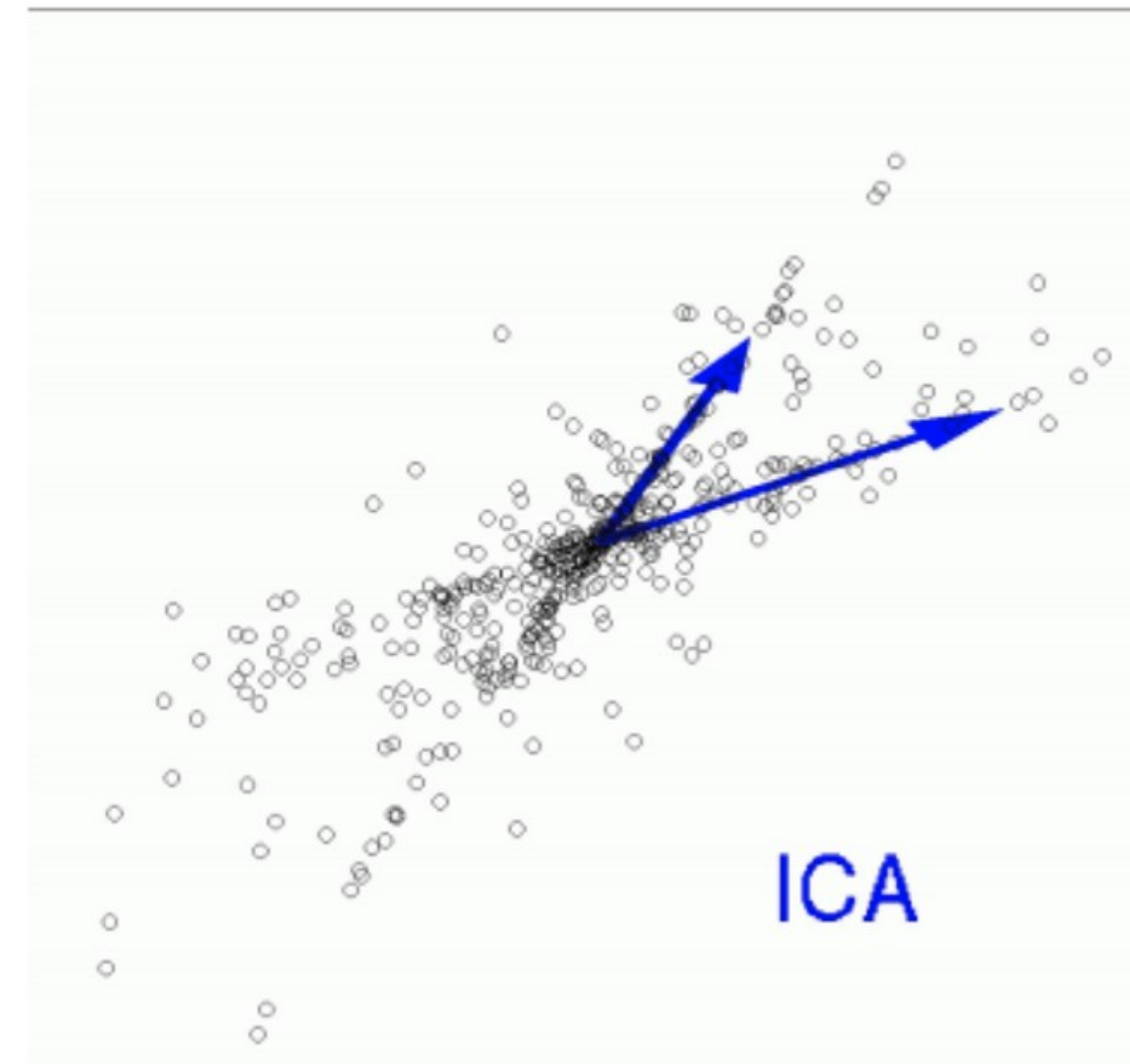
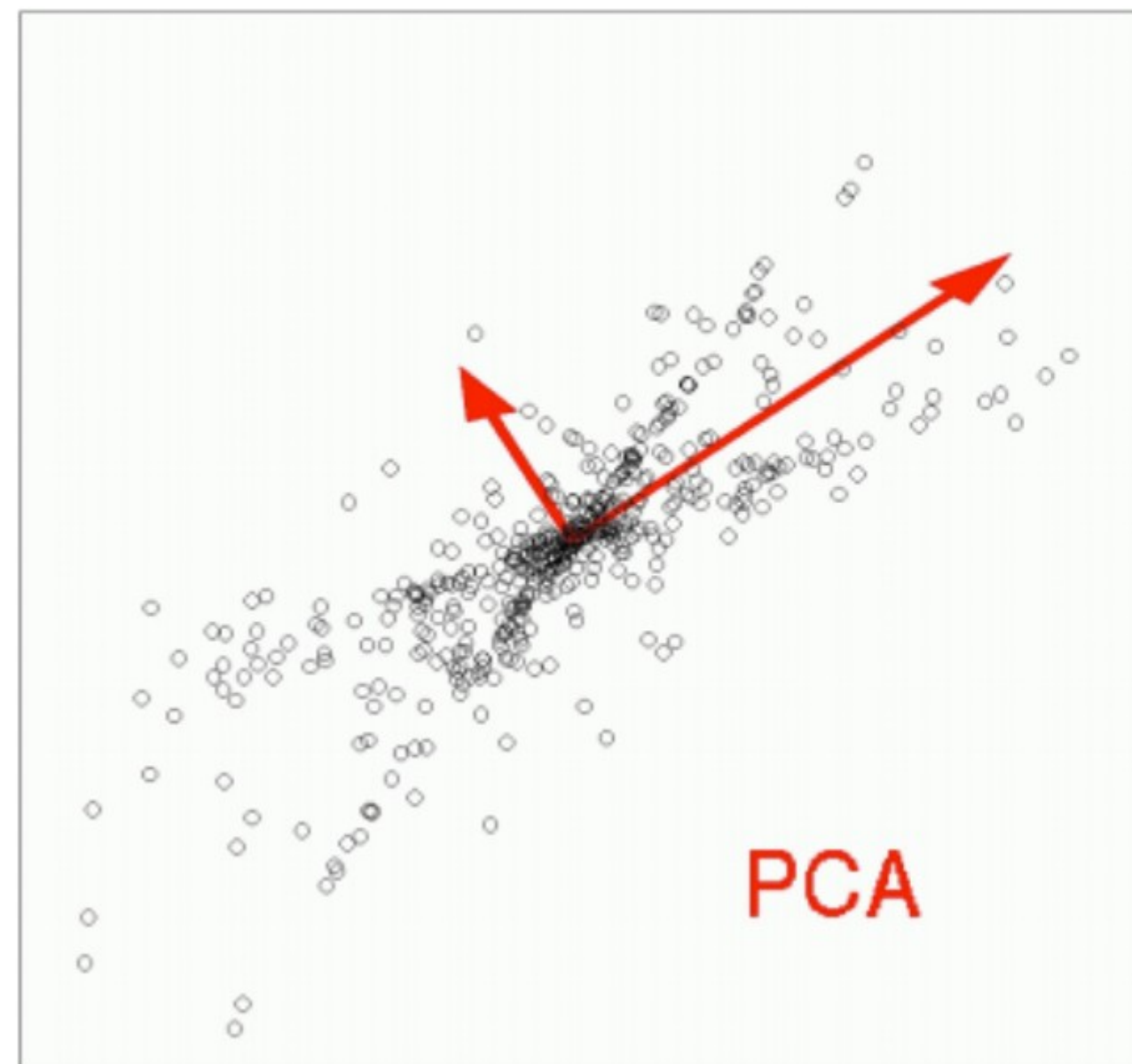
- Removes correlation
- Some components (eigenvalues) are **more important** than others
- (Eigen)vectors are **orthogonal**

## ICA:

- Removes correlation **and** higher-order dependence
- All components are **equally important**
- ICA vectors are **not orthogonal**

[http://compneurosci.com/wiki/images/4/42/Intro\\_to\\_PCA\\_and\\_ICA.pdf](http://compneurosci.com/wiki/images/4/42/Intro_to_PCA_and_ICA.pdf)

# Difference between PCA and ICA



[http://compneurosci.com/wiki/images/4/42/Intro\\_to\\_PCA\\_and\\_ICA.pdf](http://compneurosci.com/wiki/images/4/42/Intro_to_PCA_and_ICA.pdf)



Thank you for your attention!

