

Brain  
Code  
Camp

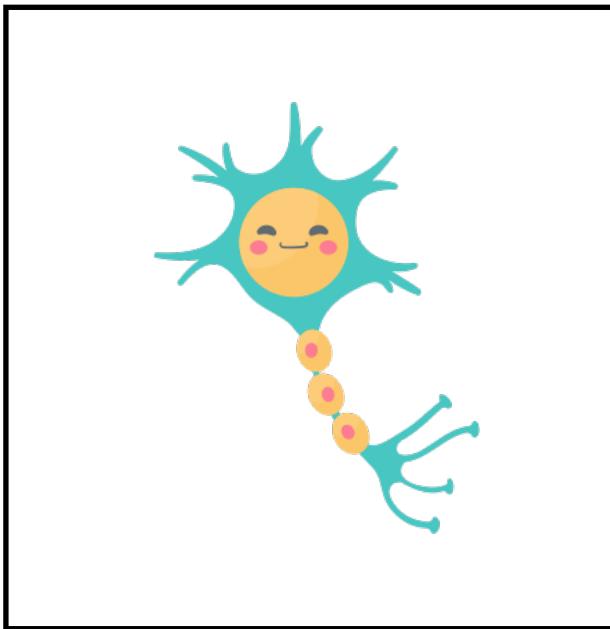
# Dimensionality Reduction

Itthi Chatnuntawech

Speaker: Itthi Chatnuntawech

Module: Dimensionality Reduction

# RGB Images



1024 x 1024 pixels

If we use raw pixels as features, we will have

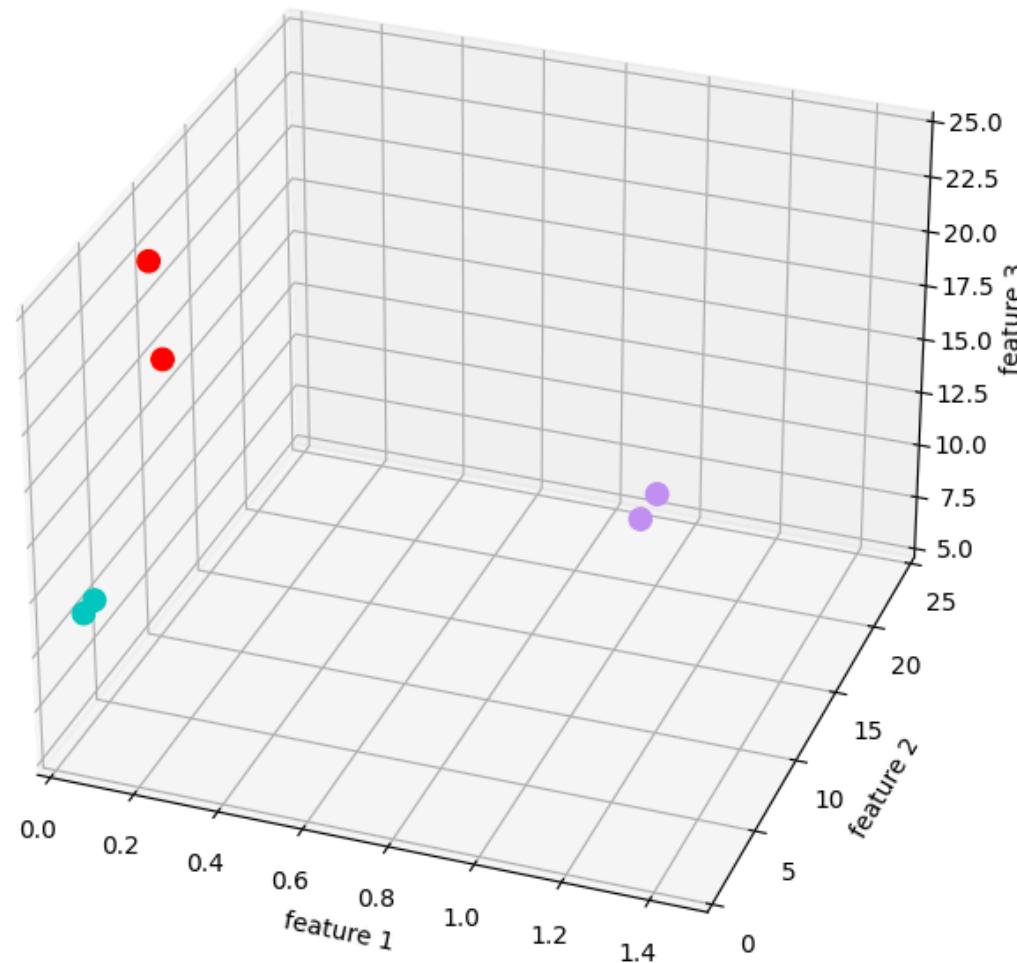
$$1024 \times 1024 \times 3 = 3,145,728 \text{ features.}$$

# of            # of            # of  
rows            columns        channels

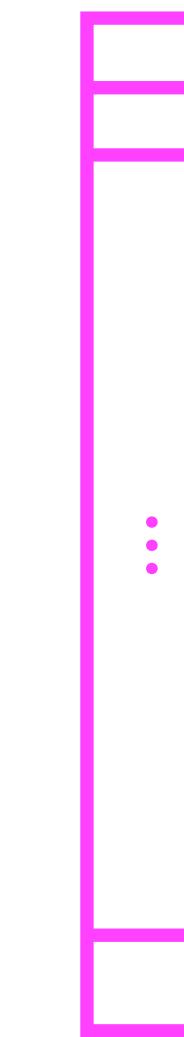
3,145,728-dimensional feature space!

# Dimensionality Reduction

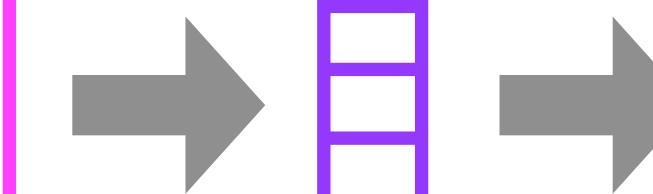
data visualization



high-dim  
space

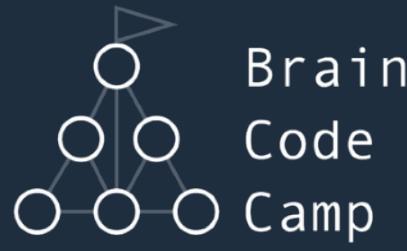


low-dim  
space



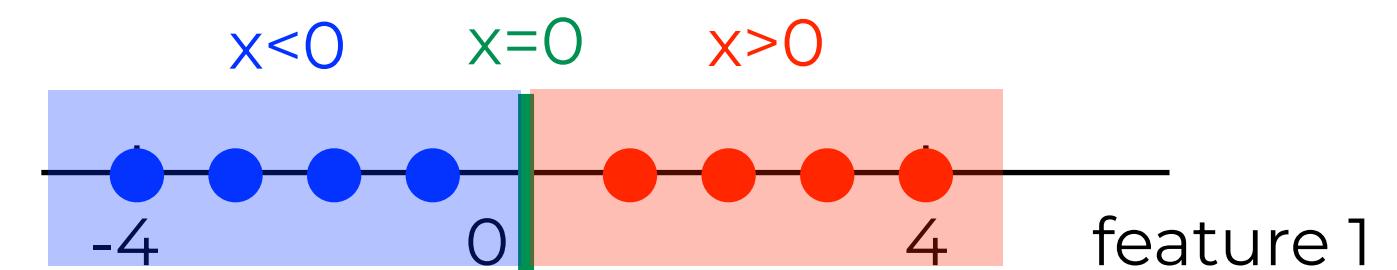
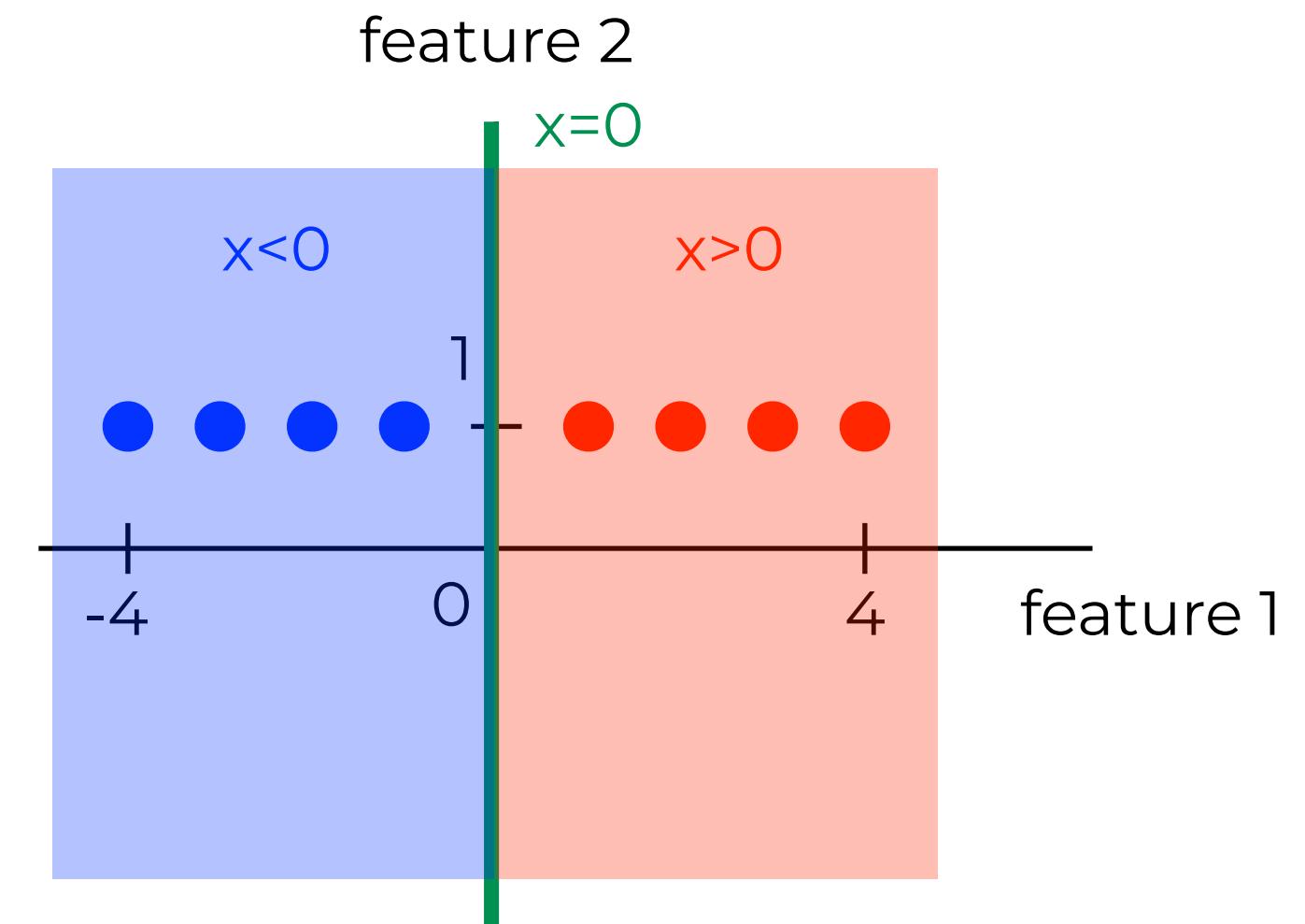
data analysis in  
lower dimensional  
space





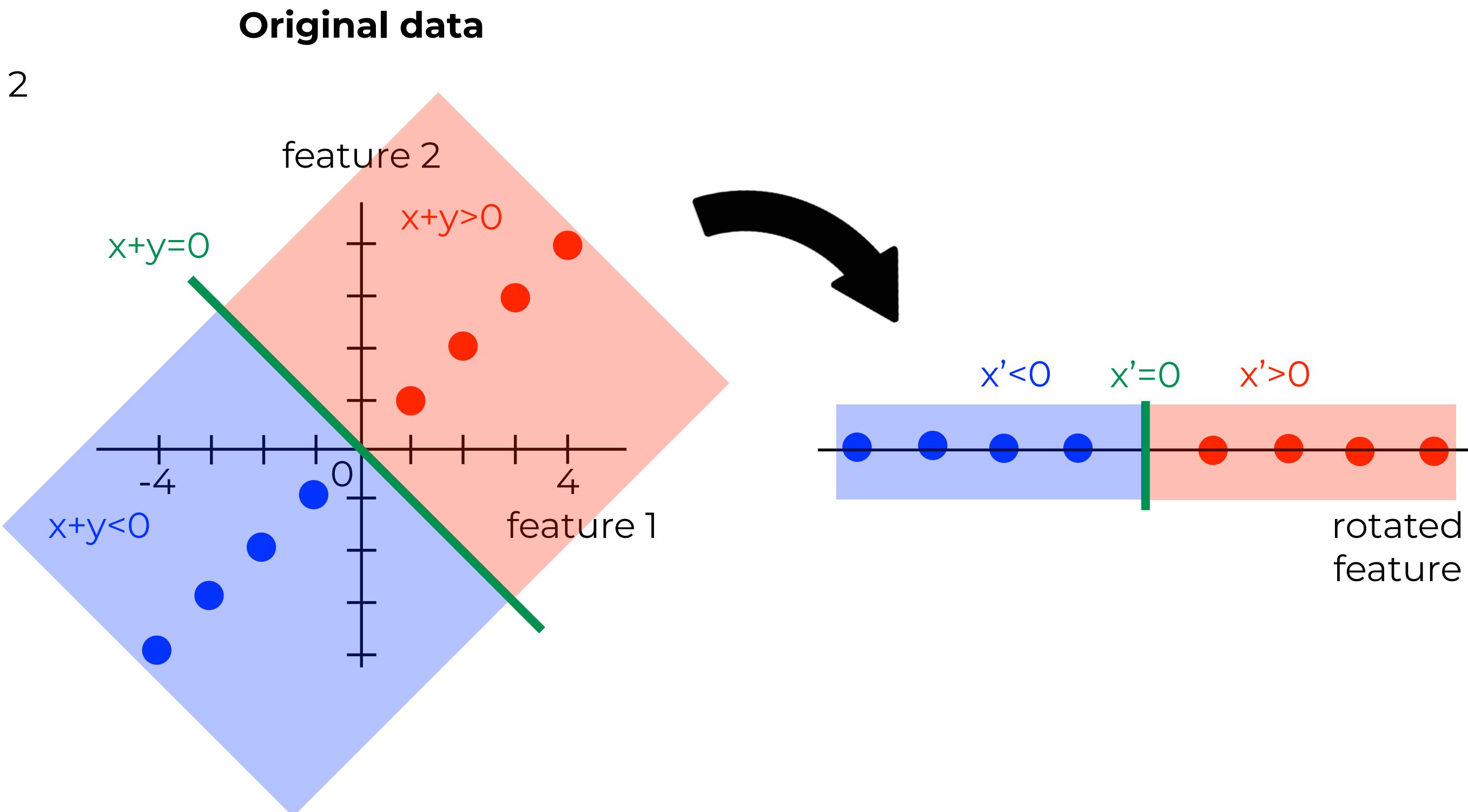
# A Simple Classification Example

|          | feature 1 | feature 2 |
|----------|-----------|-----------|
| sample 1 | -4        | 1         |
| sample 2 | -3        | 1         |
| sample 3 | -2        | 1         |
| sample 4 | -1        | 1         |
| sample 5 | 1         | 1         |
| sample 6 | 2         | 1         |
| sample 7 | 3         | 1         |
| sample 8 | 4         | 1         |



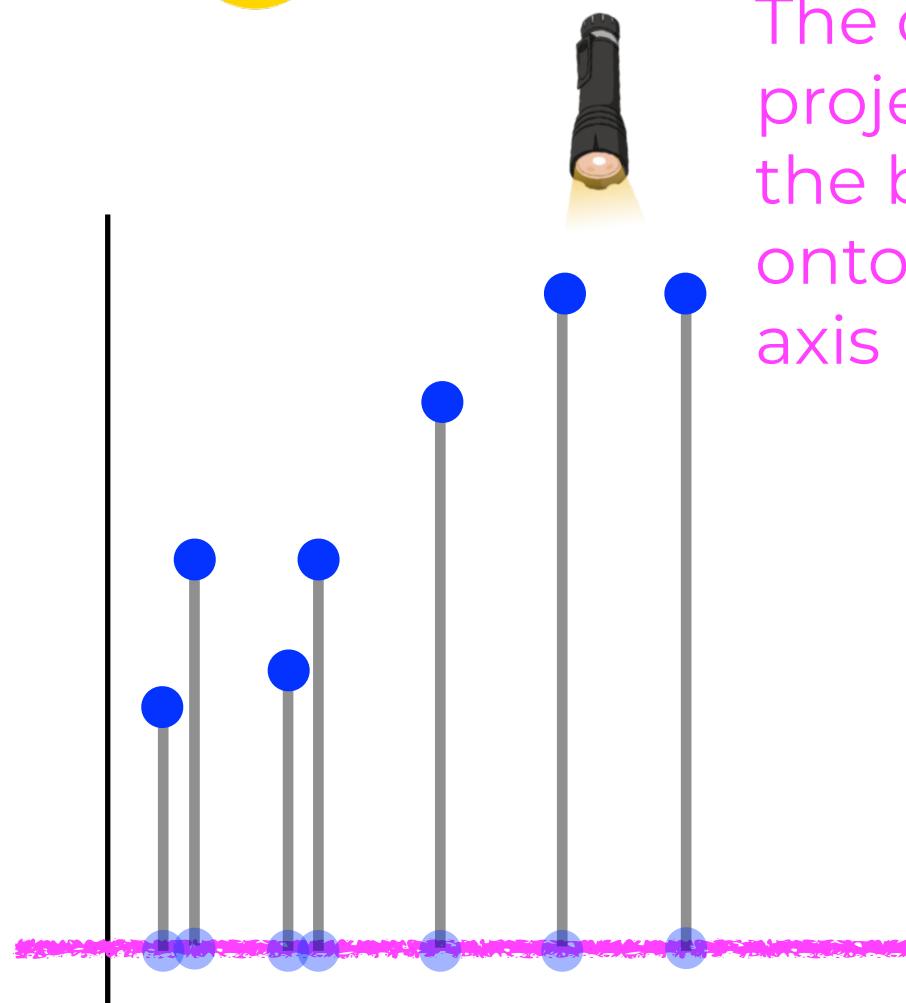
# A Simple Classification Example

|          | feature 1 | feature 2 |
|----------|-----------|-----------|
| sample 1 | -4        | -4        |
| sample 2 | -3        | -3        |
| sample 3 | -2        | -2        |
| sample 4 | -1        | -1        |
| sample 5 | 1         | 1         |
| sample 6 | 2         | 2         |
| sample 7 | 3         | 3         |
| sample 8 | 4         | 4         |

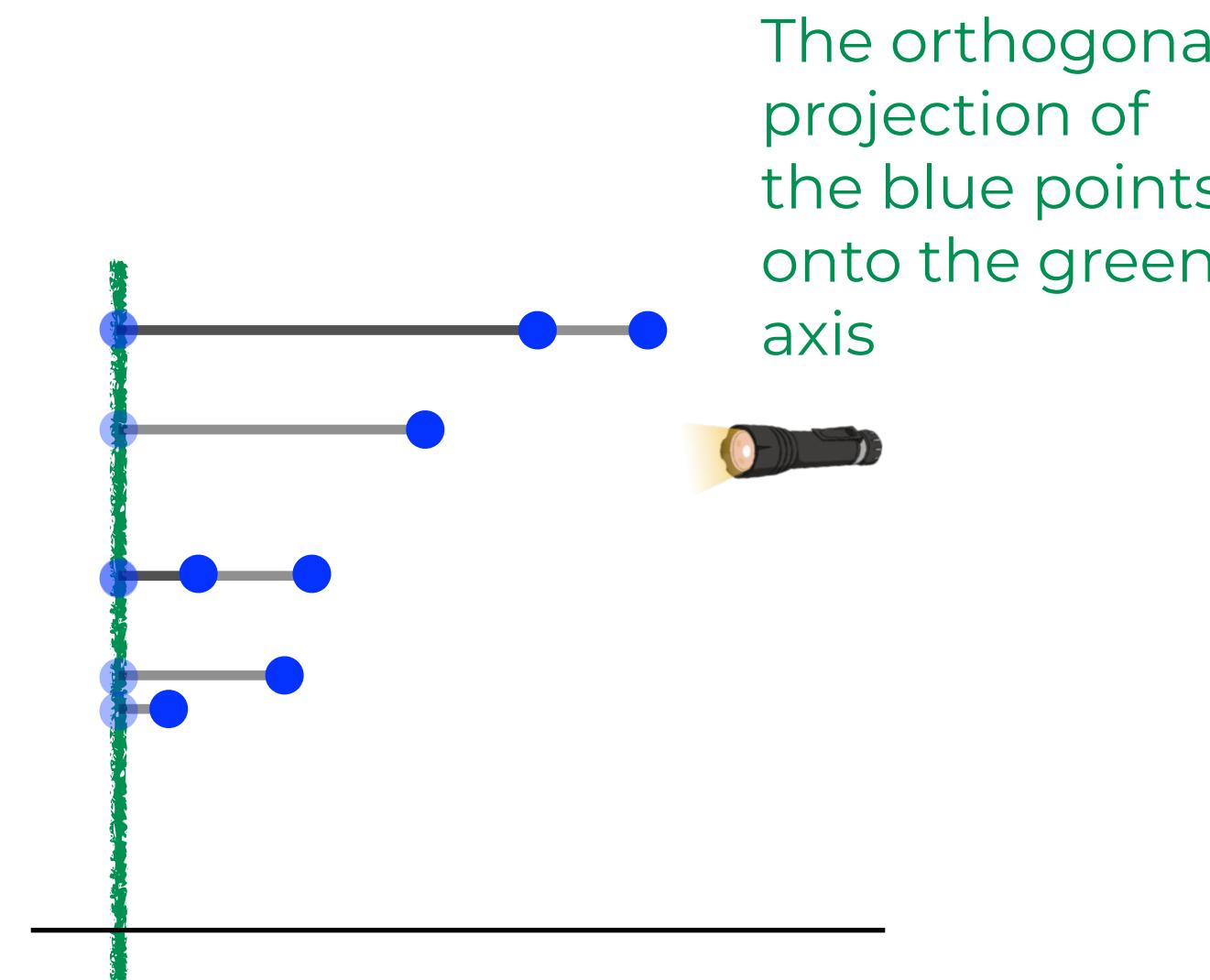


# Projection

The orthogonal projection of the blue points onto the pink axis

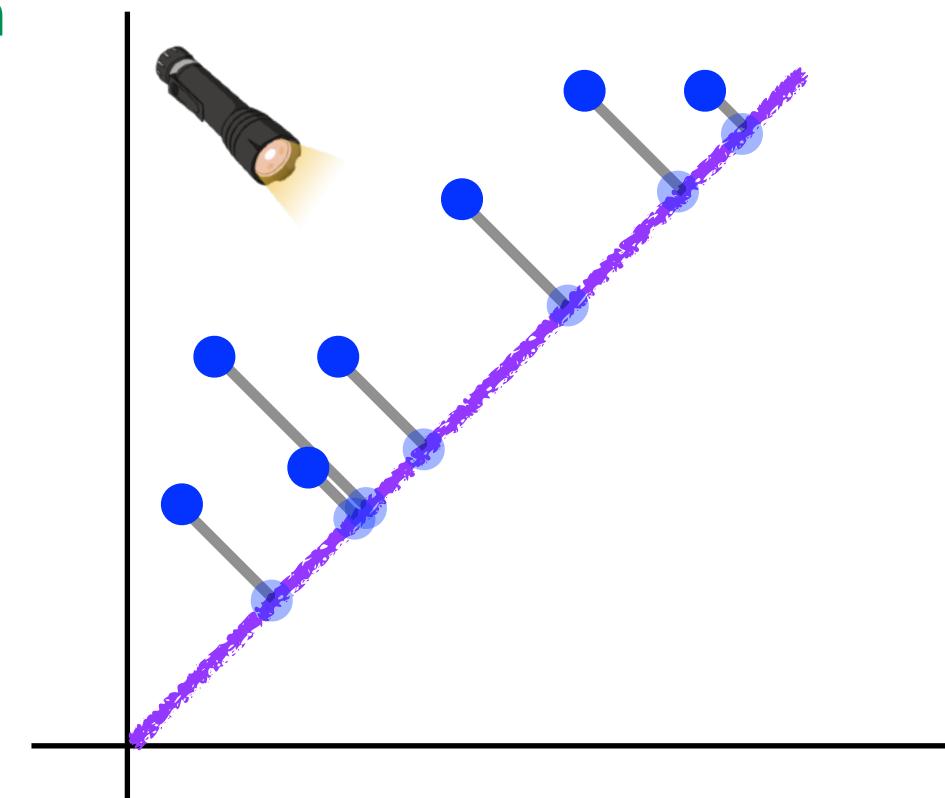


How spread out the projected points are



The orthogonal projection of the blue points onto the green axis

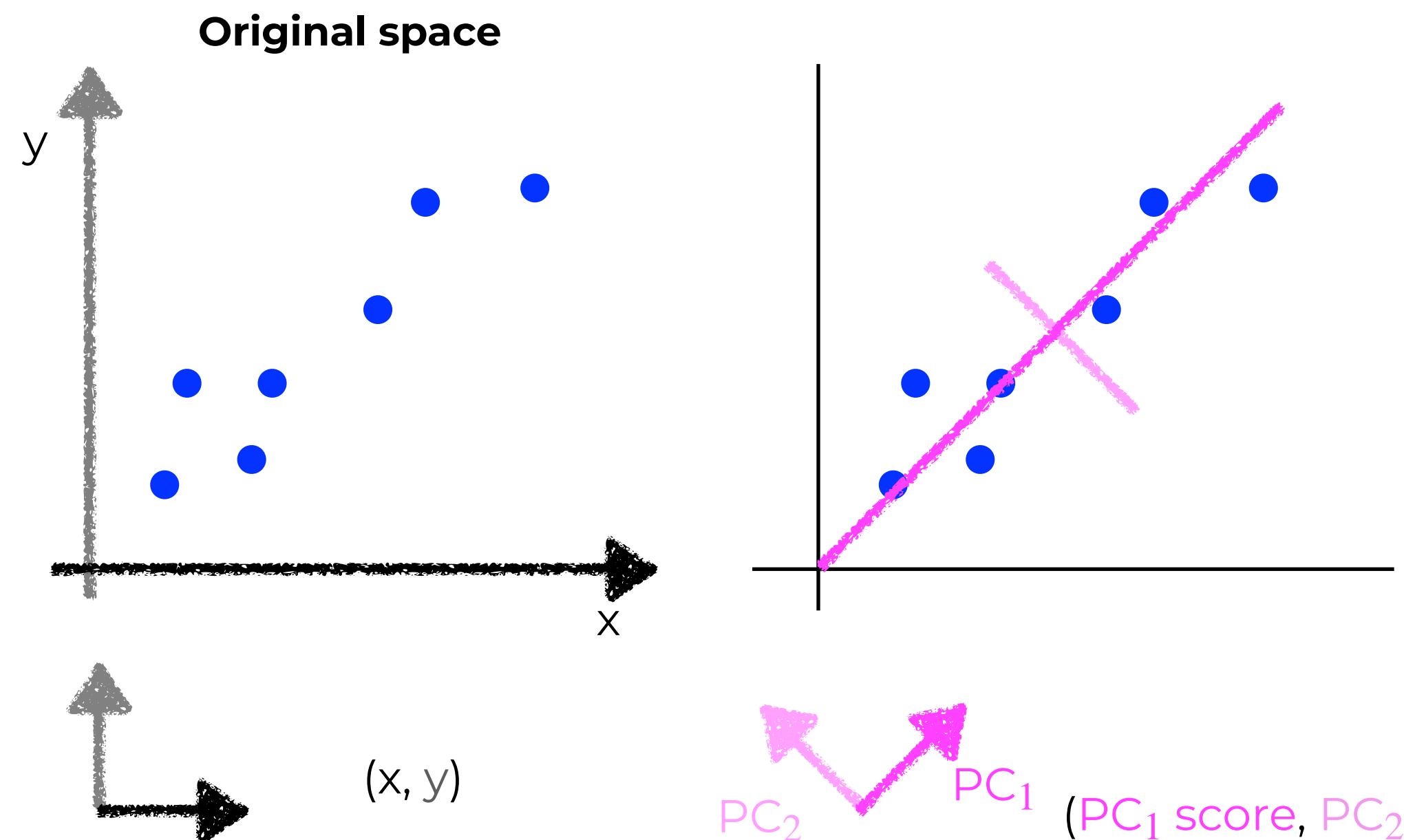
The orthogonal projection of the blue points onto the purple axis



higher

lower

# Principal Component Analysis (PCA)



For d-dimensional data, PCA can find up to d directions (principal components or PCs)

1. Find the direction that **maximizes the variance** of the projected points. This direction can be described by a vector called **the first principal component (PC<sub>1</sub>)**
2. Find the next PC that
  - is **orthogonal** to the PCs already considered
  - **maximizes the variance** along the new direction
3. Repeat step 2 until you have the desired number of PCs

# Principal Component Analysis (PCA)

## sklearn.decomposition.PCA

```
class sklearn.decomposition.PCA(n_components=None, *, copy=True, whiten=False, svd_solver='auto', tol=0.0,  
iterated_power='auto', n_oversamples=10, power_iteration_normalizer='auto', random_state=None)
```

[source]

### Attributes:

#### **components\_** : ndarray of shape (n\_components, n\_features)

Principal axes in feature space, representing the directions of maximum variance in the data. Equivalently, the right singular vectors of the centered input data, parallel to its eigenvectors. The components are sorted by decreasing `explained_variance_`.

PC<sub>1</sub>, PC<sub>2</sub>, ...

#### **explained\_variance\_** : ndarray of shape (n\_components,)

The amount of variance explained by each of the selected components. The variance estimation uses `n_samples - 1` degrees of freedom.

$\lambda_1, \lambda_2, \dots$

Equal to `n_components` largest eigenvalues of the covariance matrix of X.

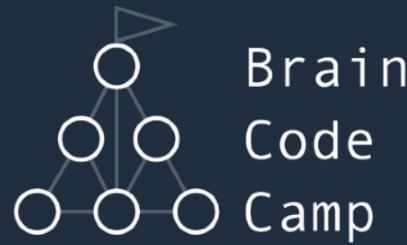
New in version 0.18.

#### **explained\_variance\_ratio\_** : ndarray of shape (n\_components,)

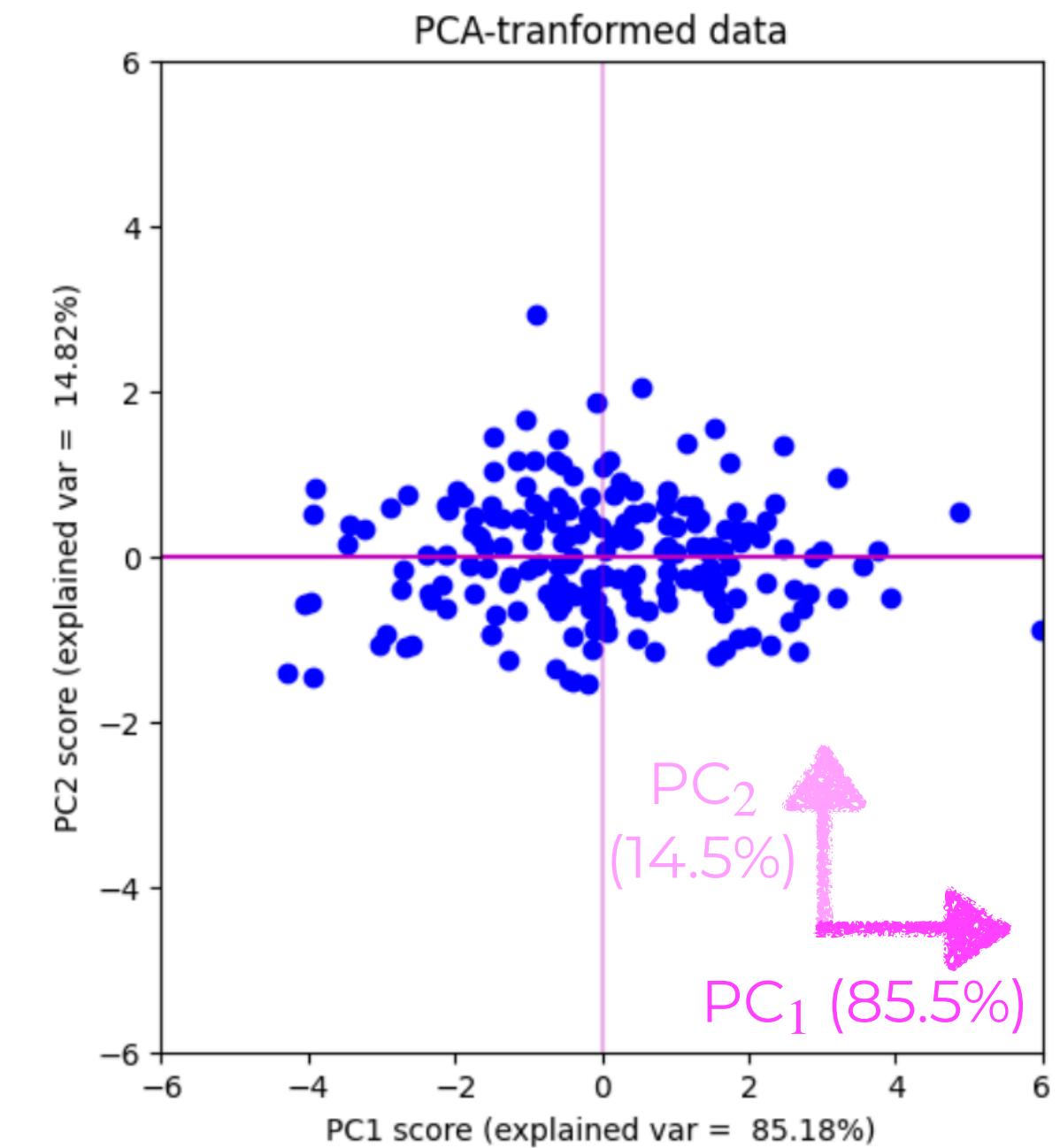
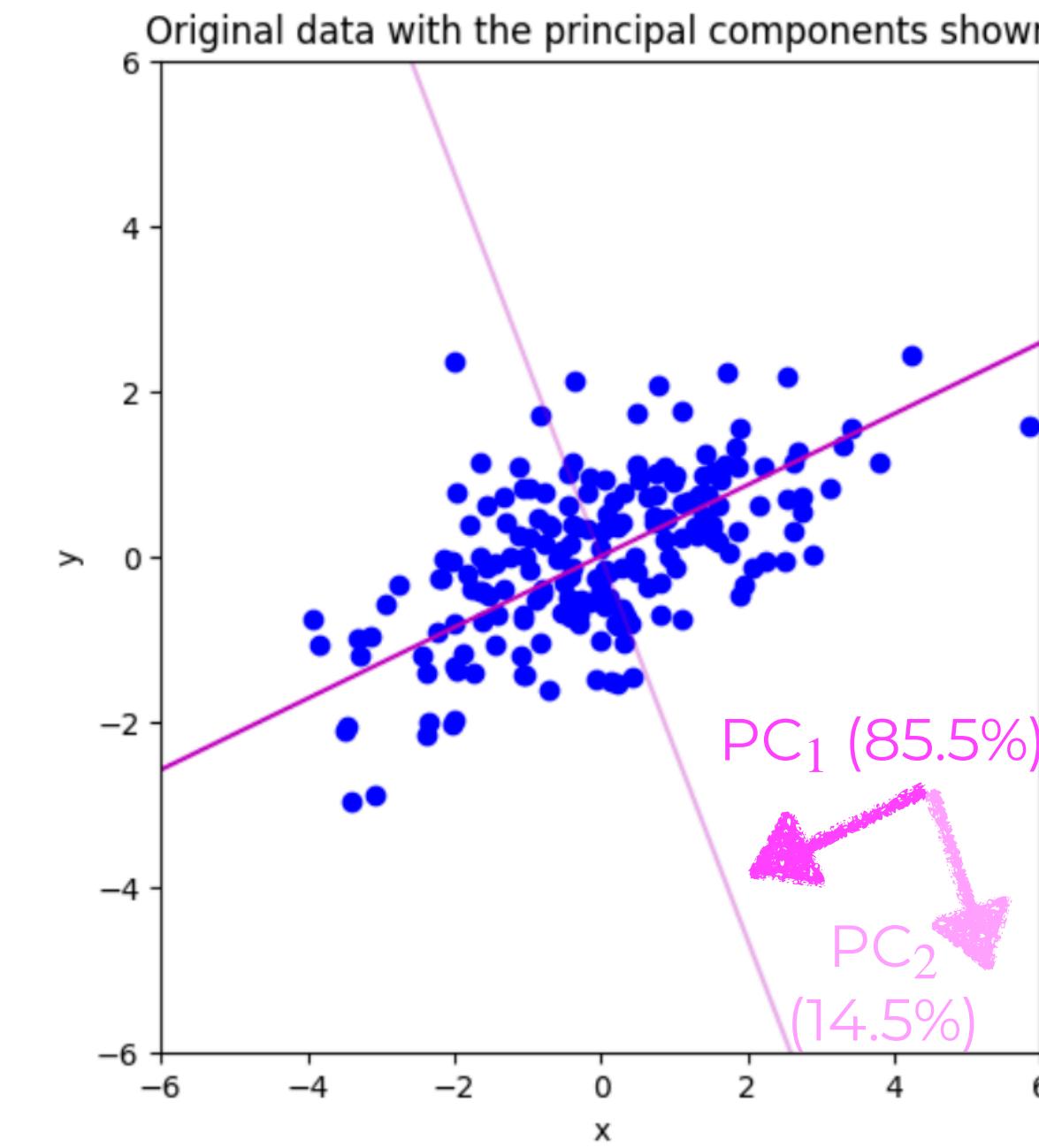
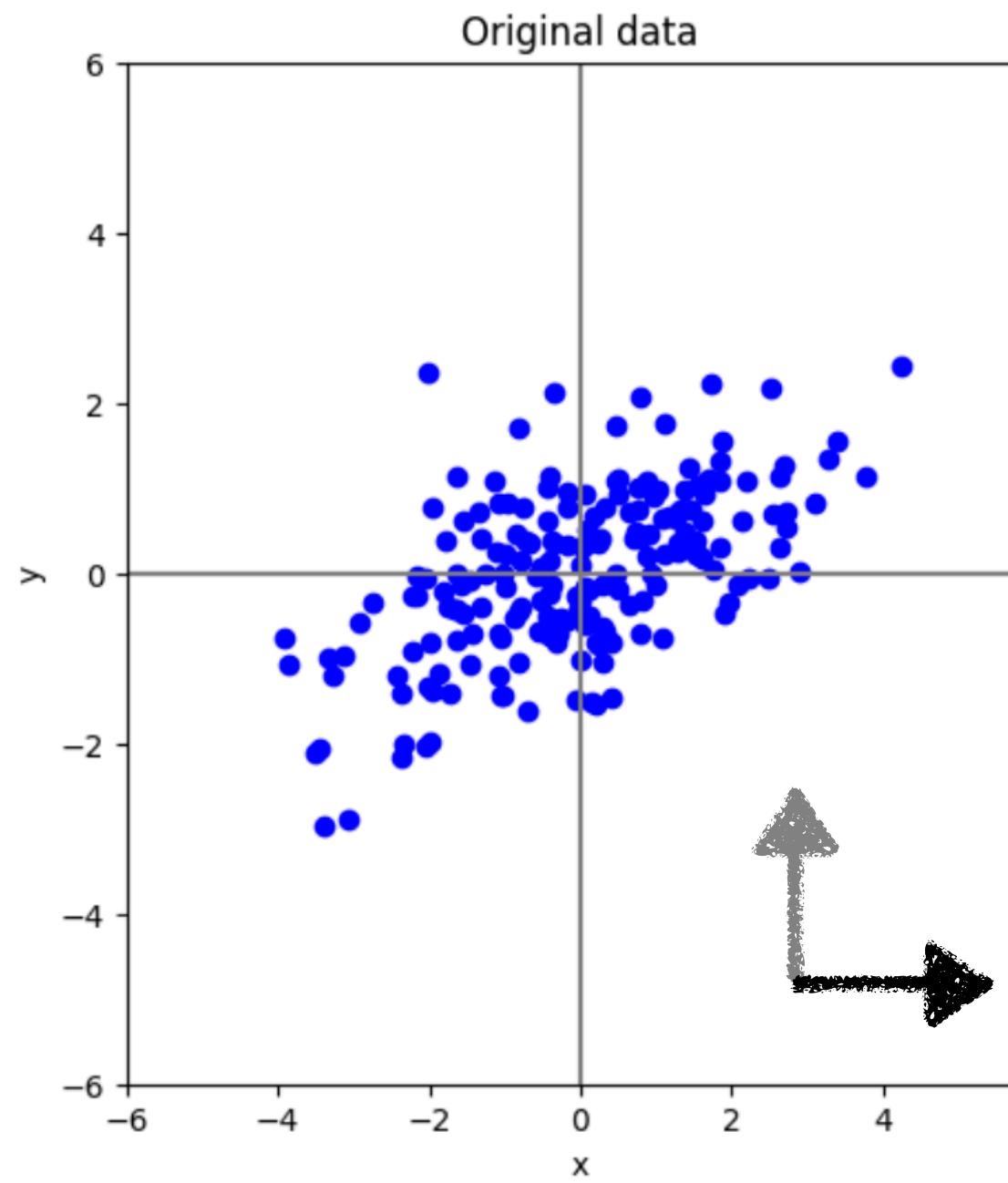
Percentage of variance explained by each of the selected components.

ex. 0.8, 0.2, ...

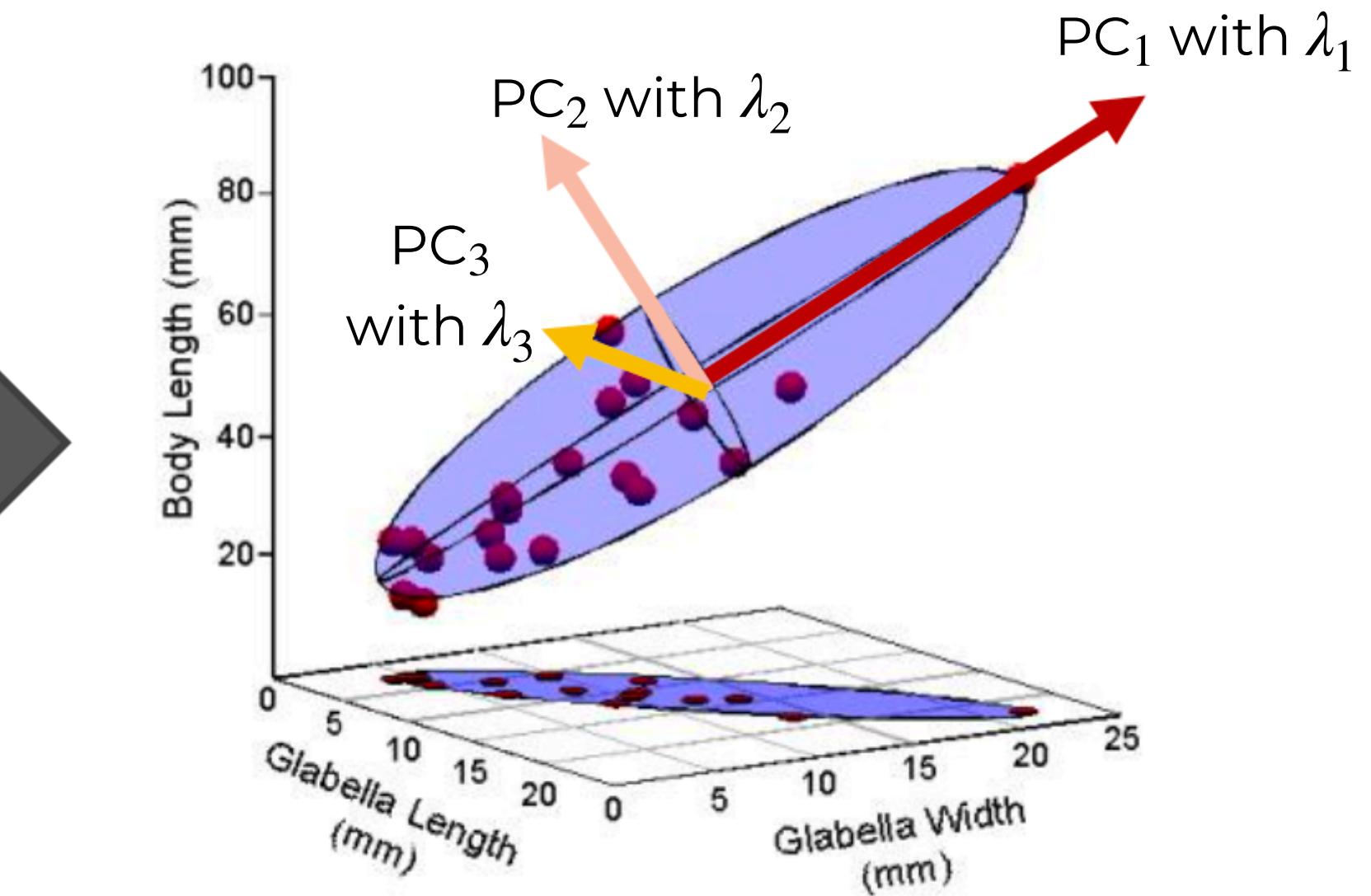
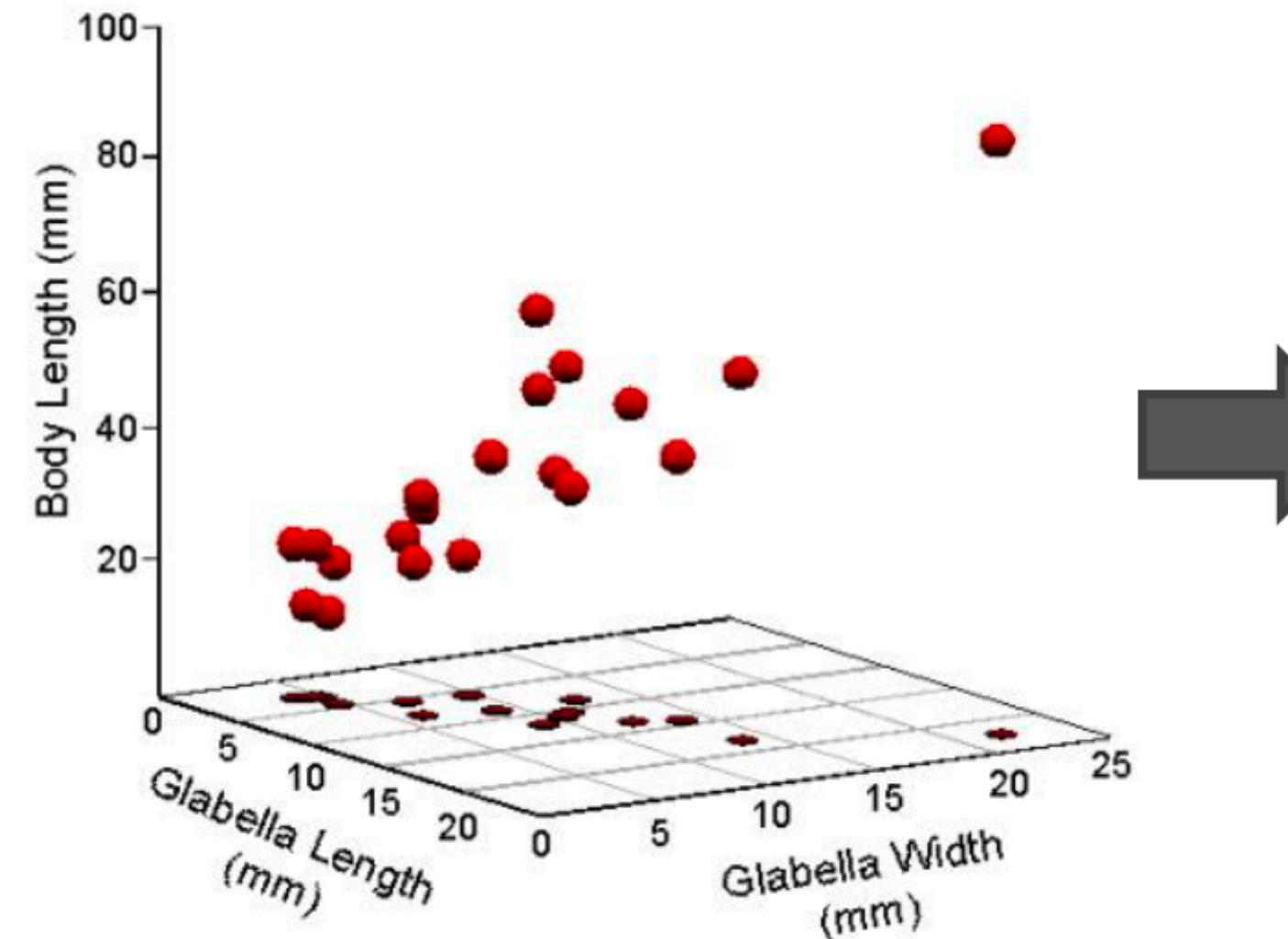
If `n_components` is not set then all components are stored and the sum of the ratios is equal to 1.0.



# PCA in 2D



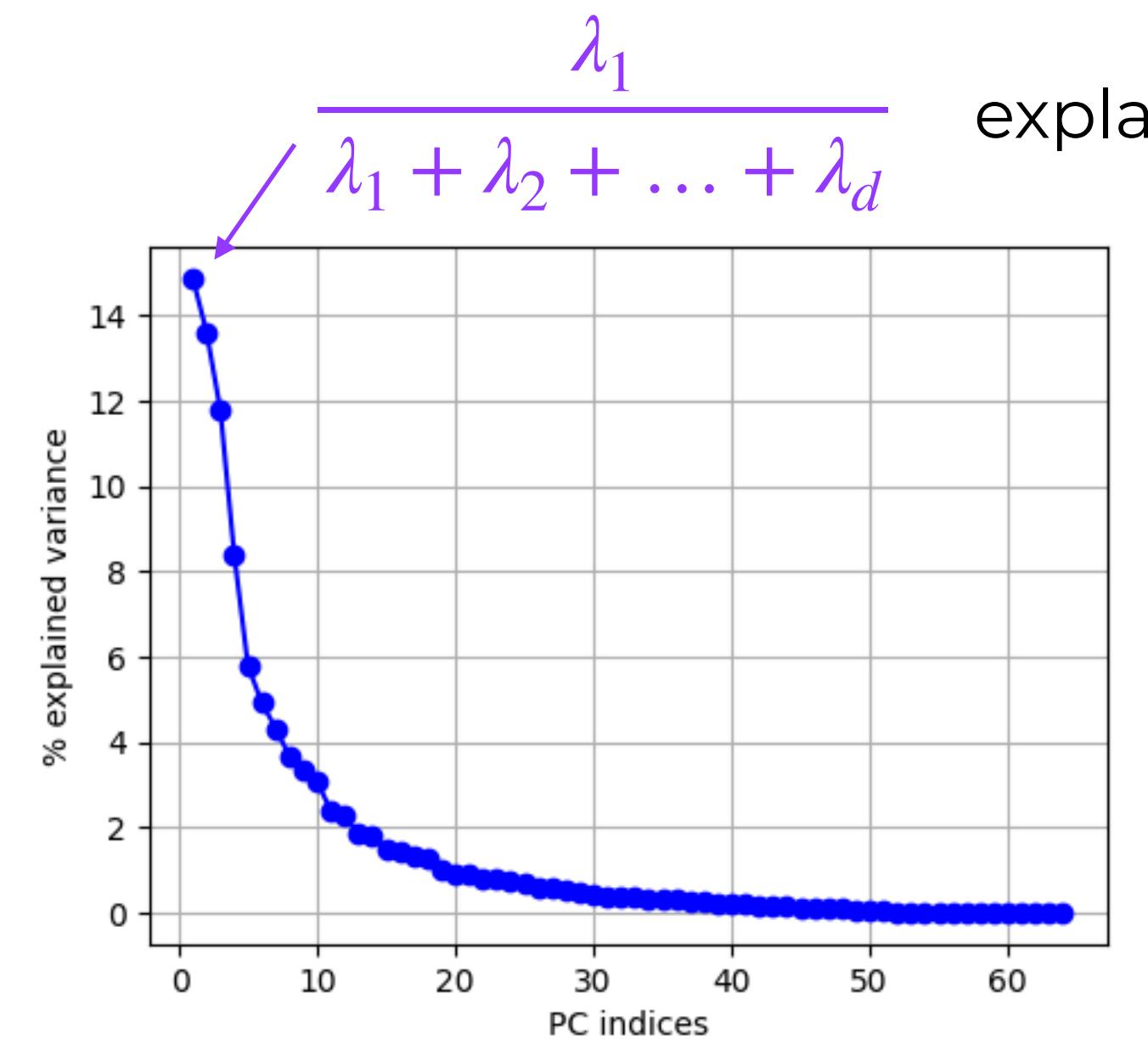
# PCA in 3D



Source: the paleontological association

# PCA in High-dimensional Space

**d-dimensional space**

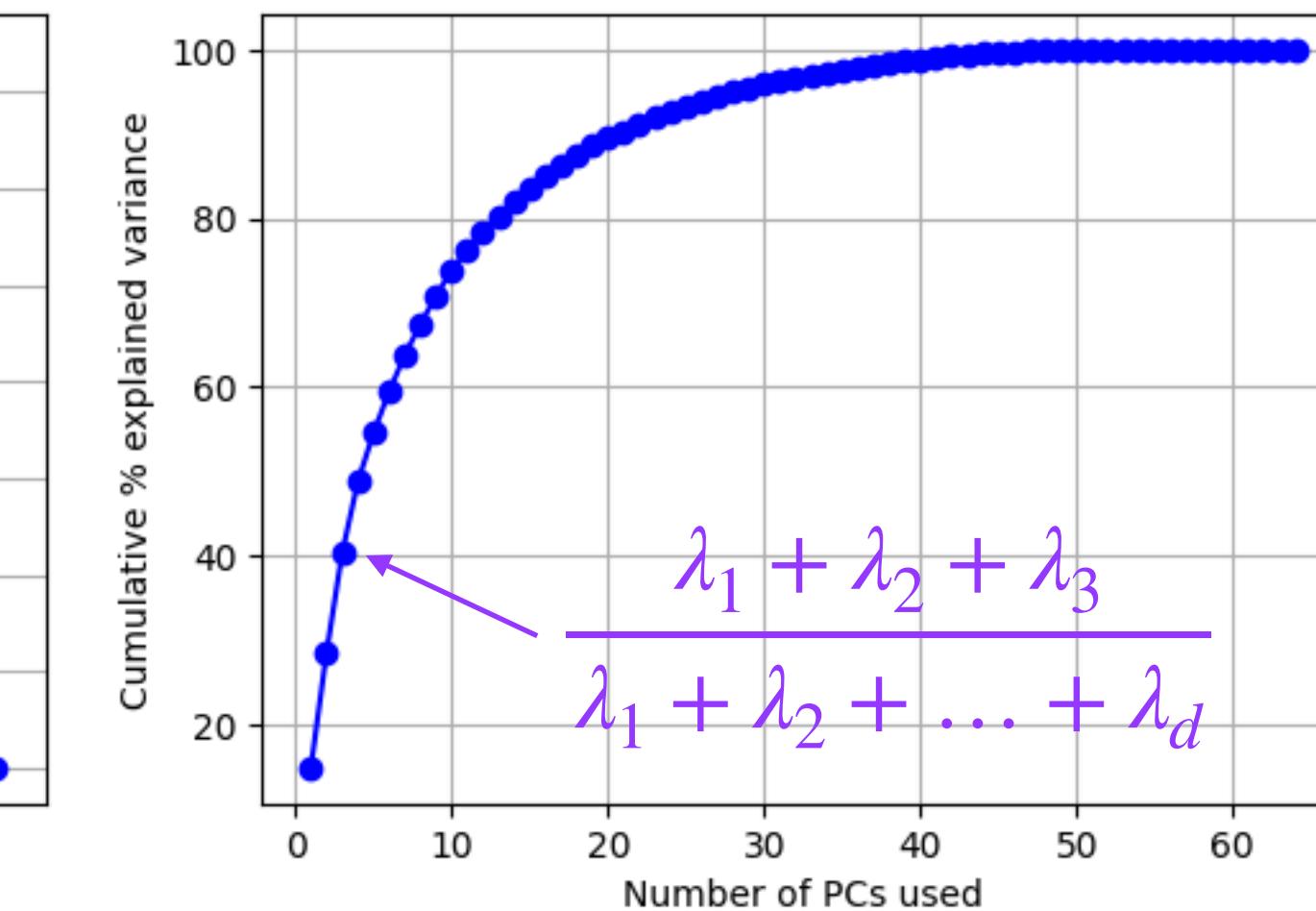


d principal components (PCs)

explained variances

$\text{PC}_1, \text{PC}_2, \dots, \text{PC}_d$

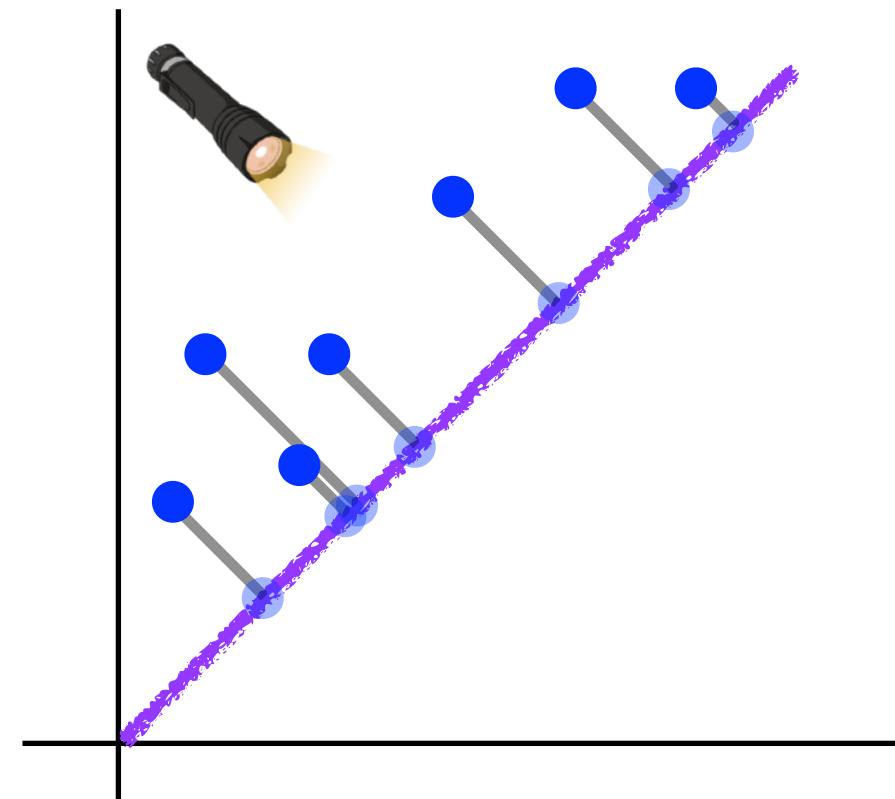
$$\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_d$$



$$\frac{\lambda_1 + \lambda_2 + \dots + \lambda_3}{\lambda_1 + \lambda_2 + \dots + \lambda_d}$$

# Two Common Definitions of PCA

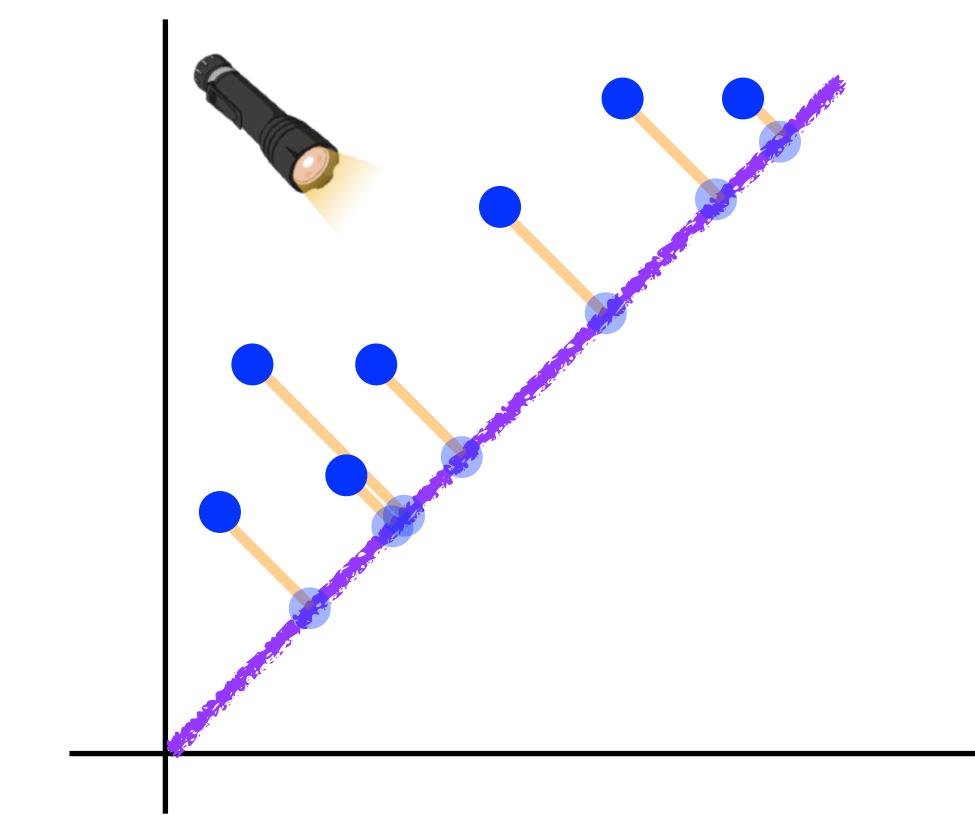
Maximum variance formulation



Maximizes the variance of the projected points

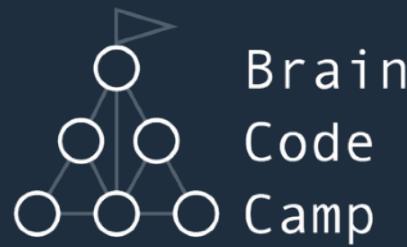


Minimum-error formulation

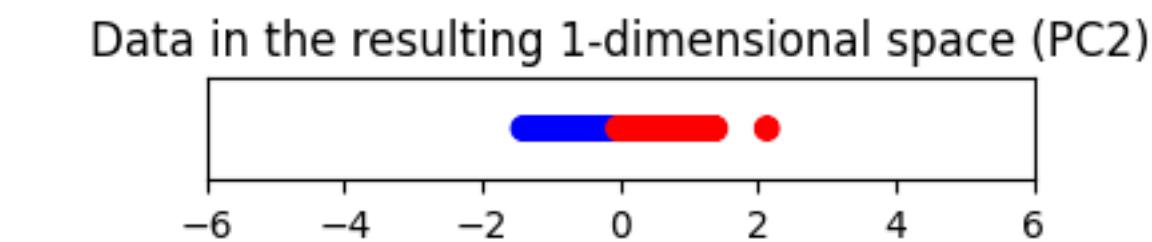
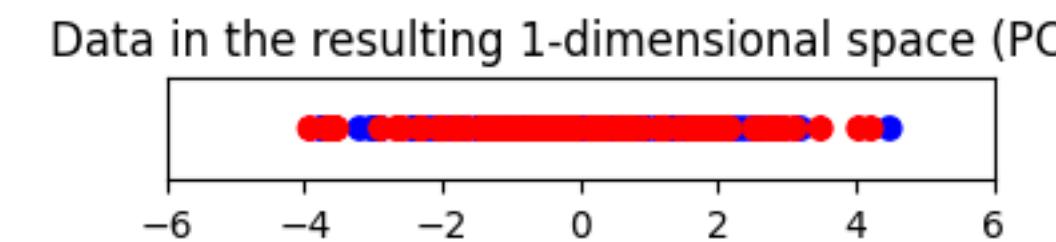
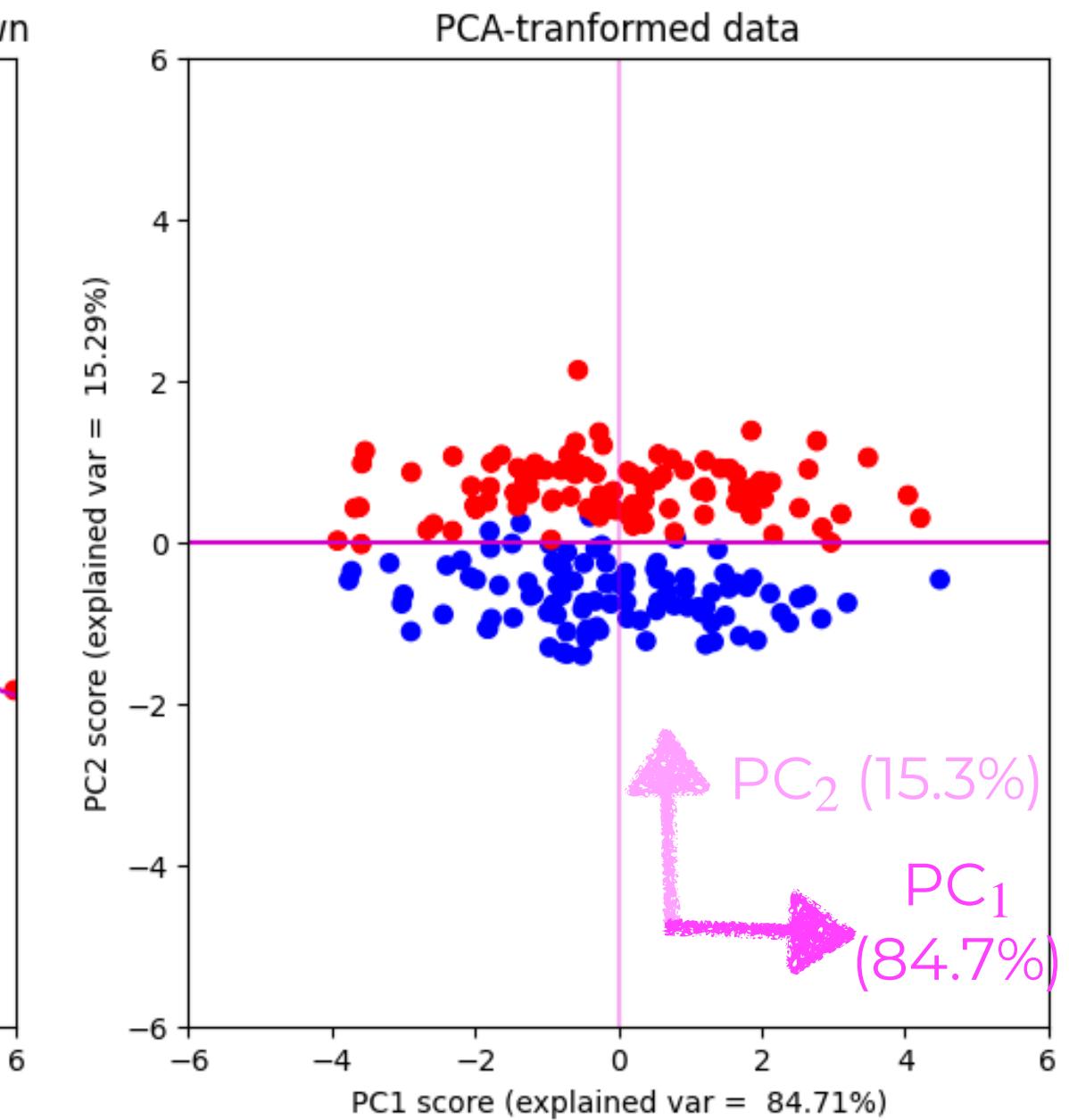
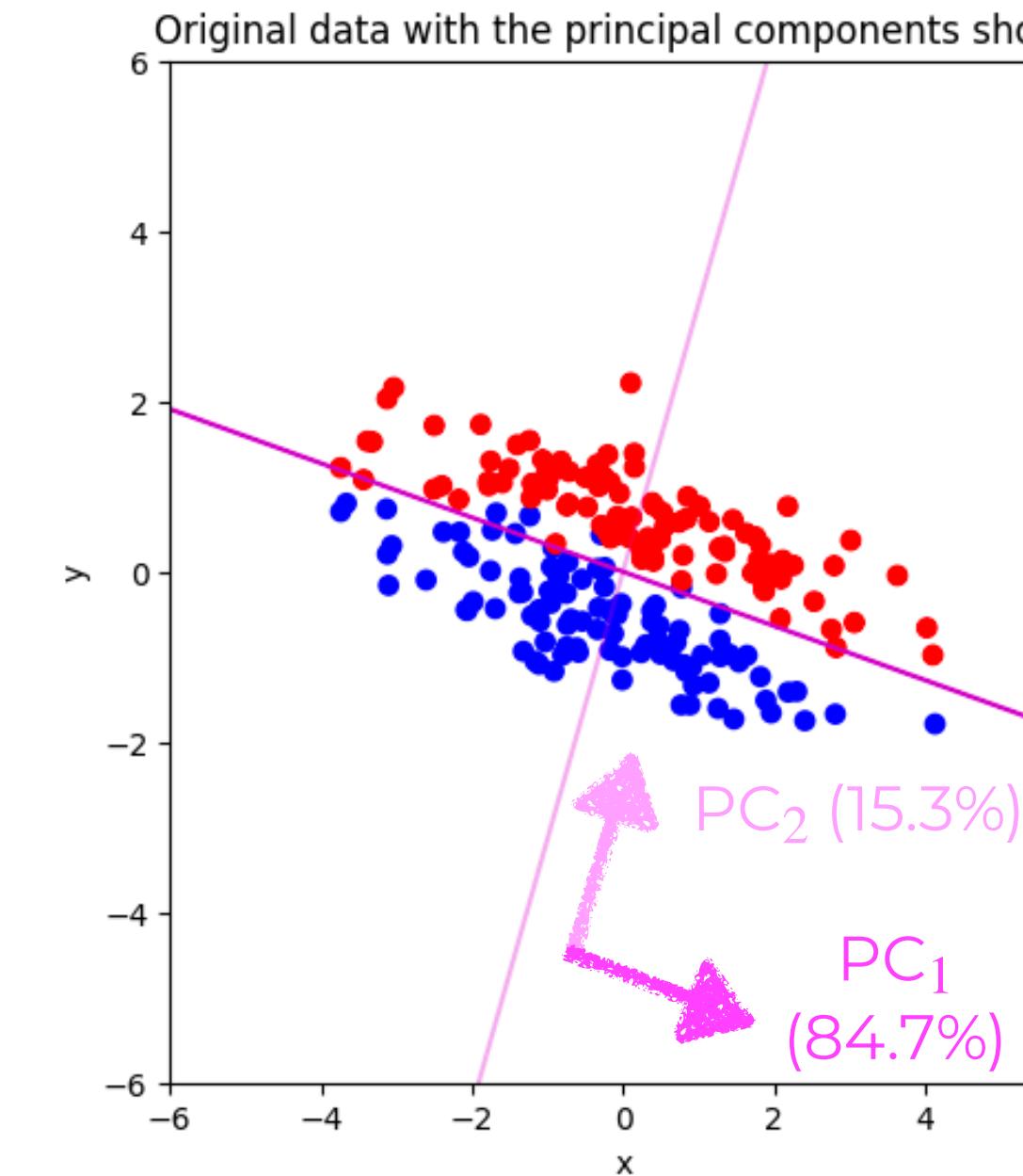
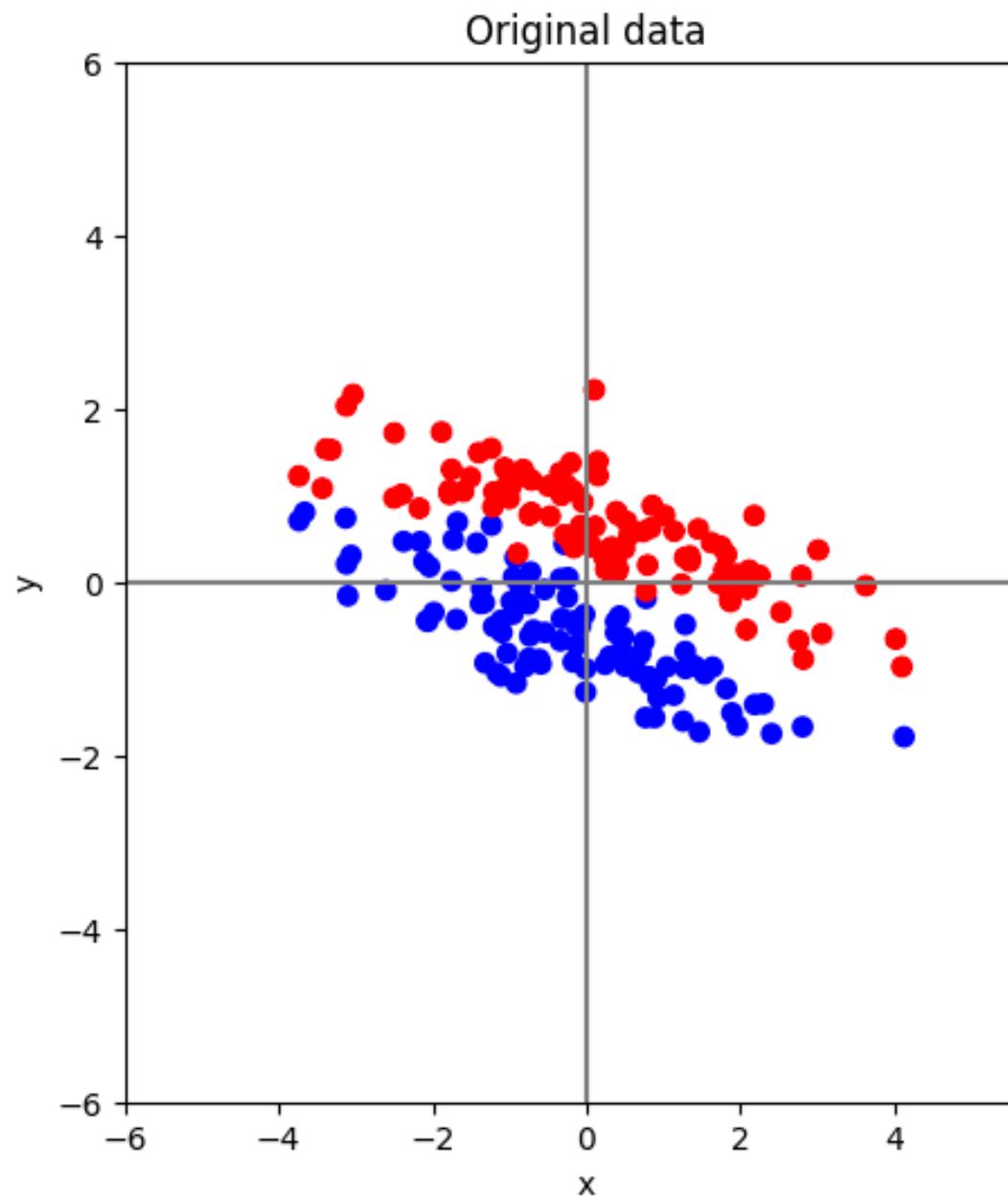


Minimizes the sum-of-squares of the projection errors

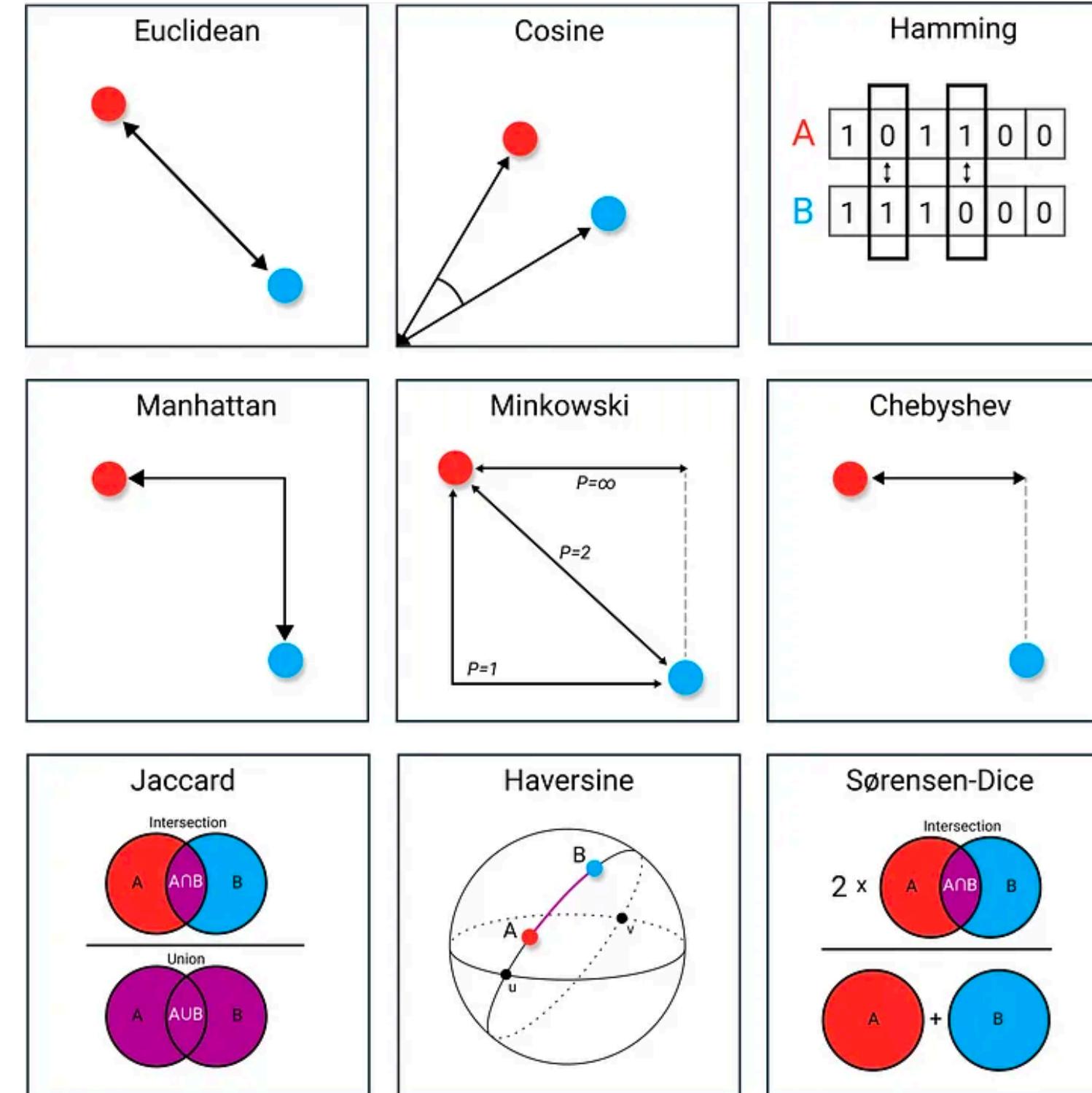
$$2 + 2 + 2 + \dots + 2 + 2$$



# PCA Example

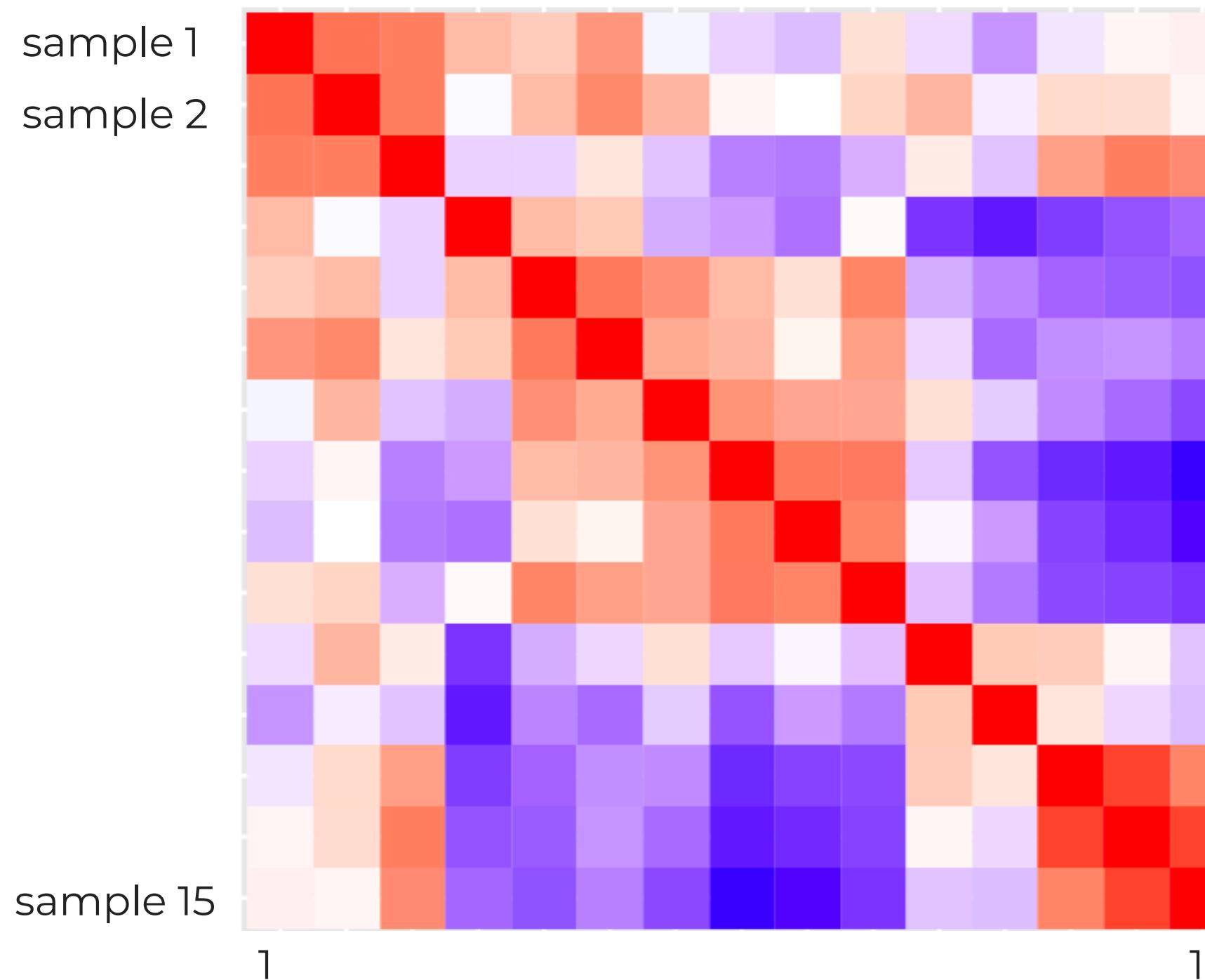


# Distance Measures



<https://towardsdatascience.com/9-distance-measures-in-data-science-918109d069fa>

# Pairwise Distance Matrix

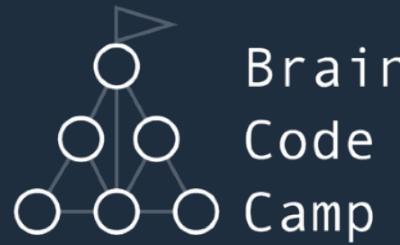


The element at row  $i$  and column  $j$  represents the distance between sample  $i$  and sample  $j$

$$D_{ij} = D_{ji}$$

$$D_{ii} = 0$$

Modified from <https://www.datanovia.com/en/lessons/clustering-distance-measures/>

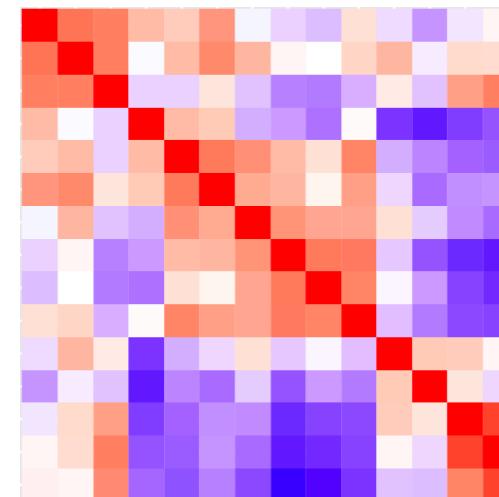


# Generalized Multi-Dimensional Scaling (MDS)

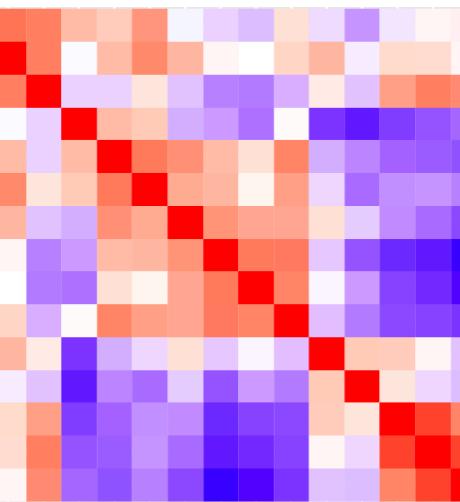
## High-dimensional feature space



pairwise distance matrix  
created using any types  
of distance



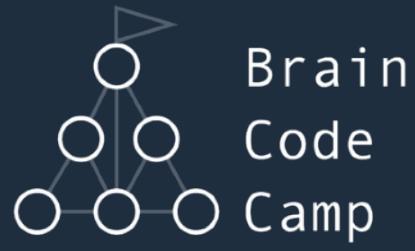
$\approx$



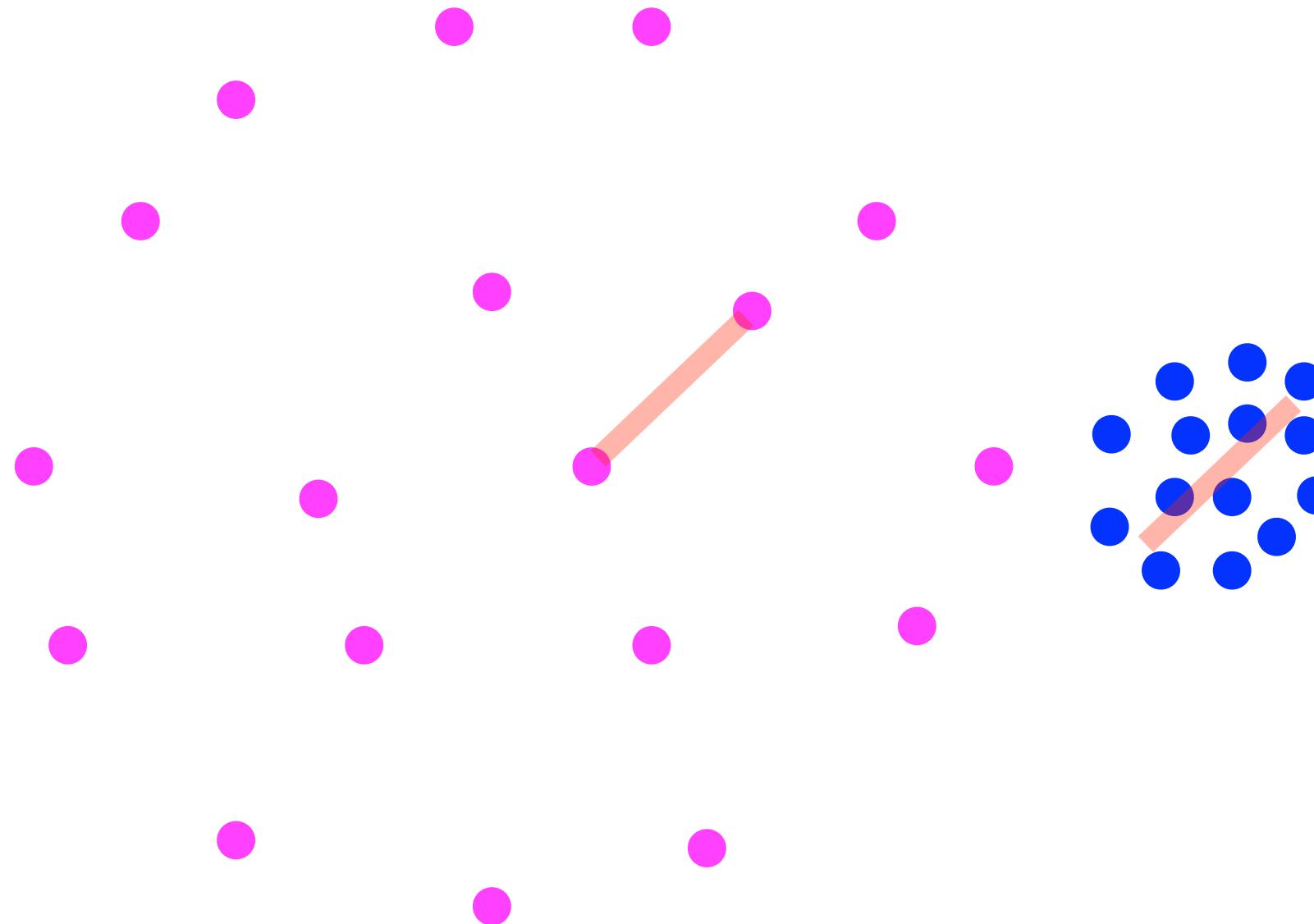
pairwise distance matrix  
created using any types  
of distance

## Low-dimensional space





# Beyond MDS



Should we use distance metrics that take into account data density?

# Popular Nonlinear Dimensionality Reduction Methods

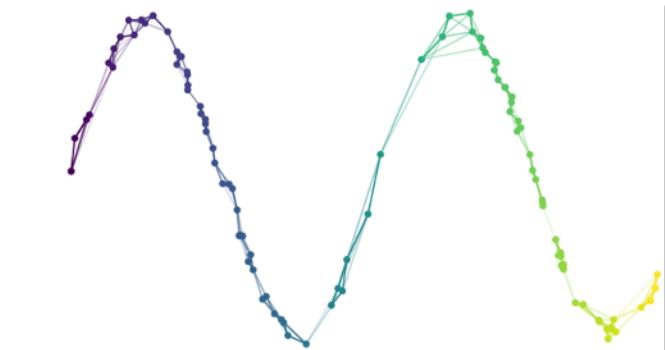
## t-distributed Stochastic Neighbor Embedding (t-SNE)

### sklearn.manifold.TSNE

```
class sklearn.manifold.TSNE(n_components=2, *, perplexity=30.0, early_exaggeration=12.0, learning_rate='auto',  
n_iter=1000, n_iter_without_progress=300, min_grad_norm=1e-07, metric='euclidean', metric_params=None, init='pca',  
verbose=0, random_state=None, method='barnes_hut', angle=0.5, n_jobs=None) [source]
```

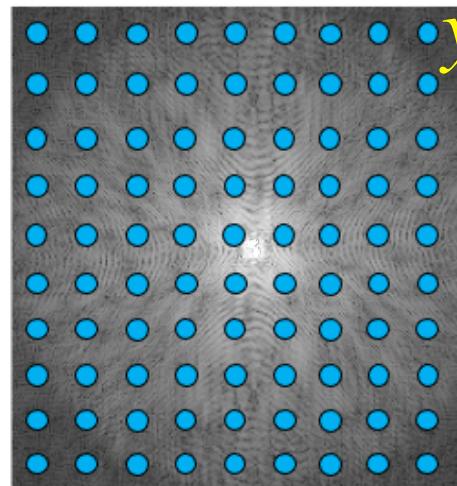
## Uniform Manifold Approximation and Projection (UMAP)

```
class umap.umap_.UMAP(n_neighbors=15, n_components=2, metric='euclidean', metric_kwds=None,  
output_metric='euclidean', output_metric_kwds=None, n_epochs=None, learning_rate=1.0, init='spectral',  
min_dist=0.1, spread=1.0, low_memory=True, n_jobs=-1, set_op_mix_ratio=1.0, local_connectivity=1.0,  
repulsion_strength=1.0, negative_sample_rate=5, transform_queue_size=4.0, a=None, b=None,  
random_state=None, angular_rp_forest=False, target_n_neighbors=-1, target_metric='categorical',  
target_metric_kwds=None, target_weight=0.5, transform_seed=42, transform_mode='embedding',  
force_approximation_algorithm=False, verbose=False, unique=False, densmap=False, dens_lambda=2.0,  
dens_frac=0.3, dens_var_shift=0.1, output_dens=False, disconnection_distance=None) [source]
```

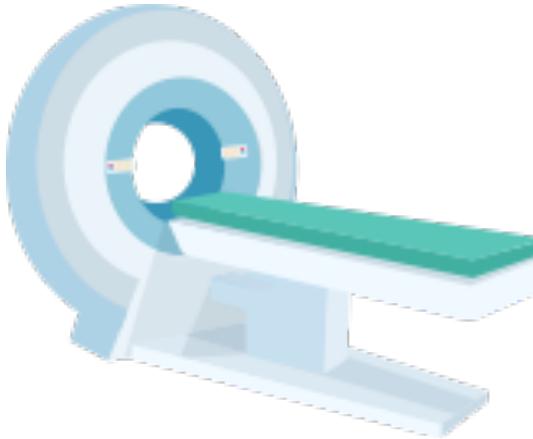


# Single-Channel MRI

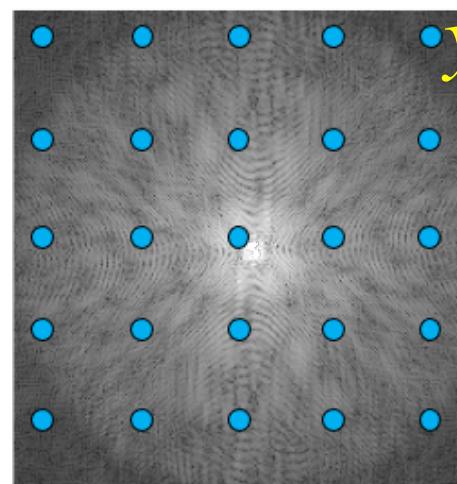
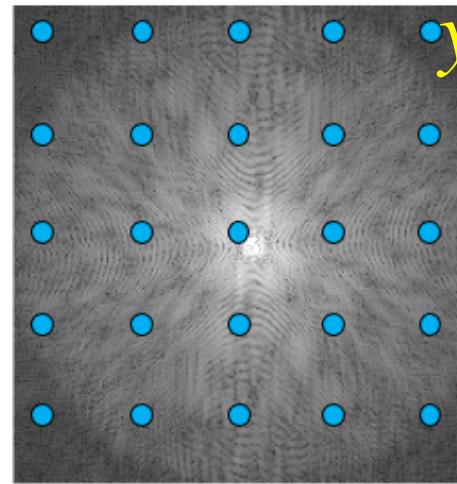
Acquired k-space data



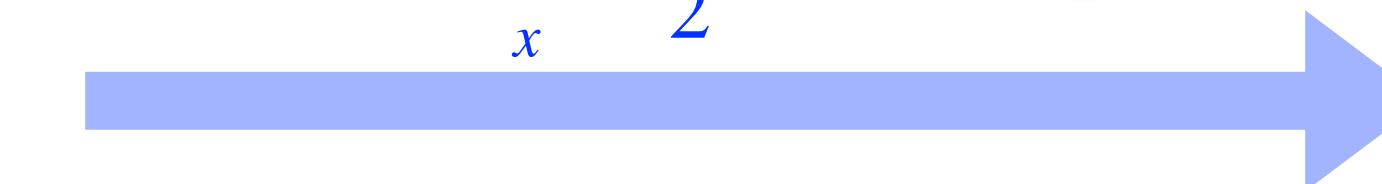
*A*



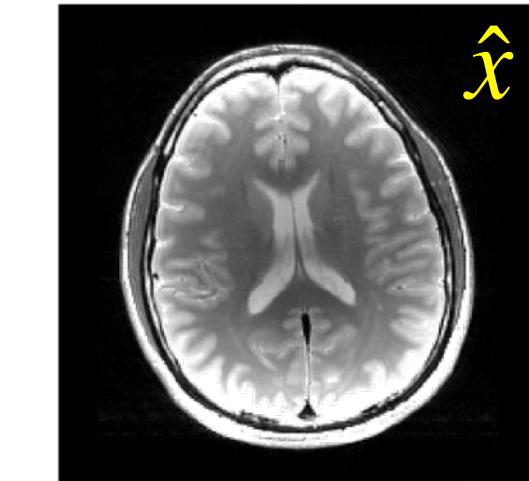
MR acquisition  
approximated using  
a linear operator



$$\hat{x} = \arg \min_x \frac{1}{2} \|Ax - y\|_2^2$$



Reconstructed data



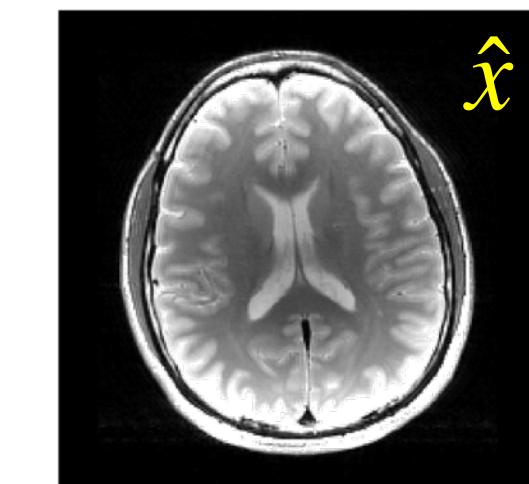
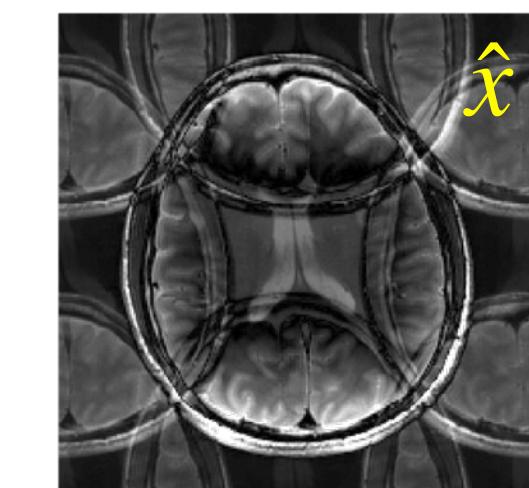
$$\hat{x} = \arg \min_x \frac{1}{2} \|Ax - y\|_2^2$$

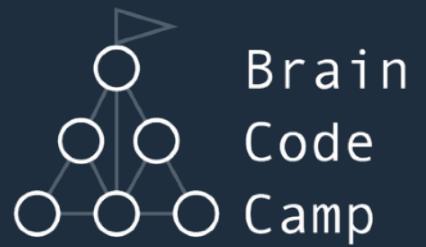


$$\hat{x} = \arg \min_x \frac{1}{2} \|Ax - y\|_2^2 + \lambda R(x)$$



Longer reconstruction time

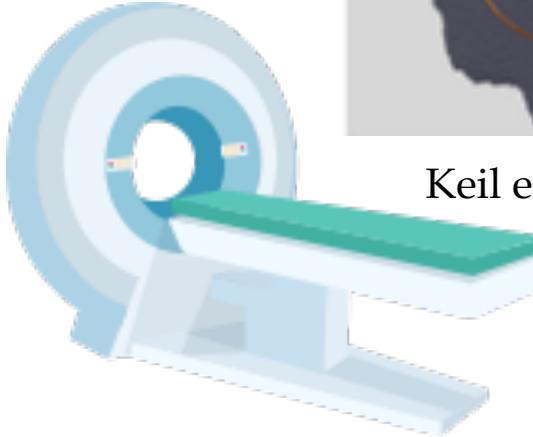




Brain  
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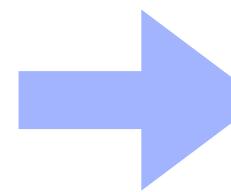
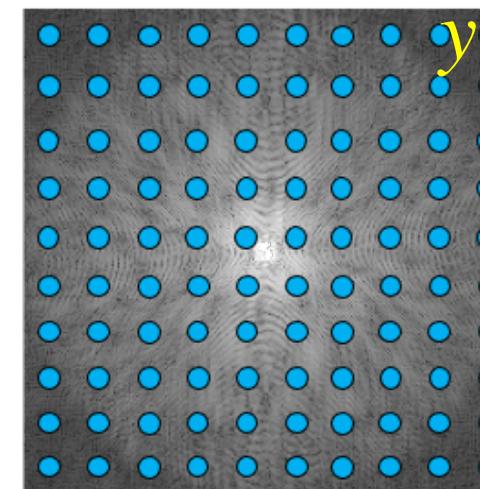
Keil et al. JMR (2013)



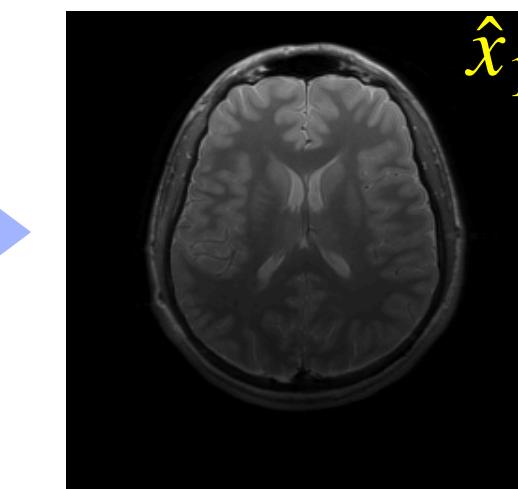
MR acquisition  
approximated using  
a linear operator

# Multi-Channel MRI

channel 1



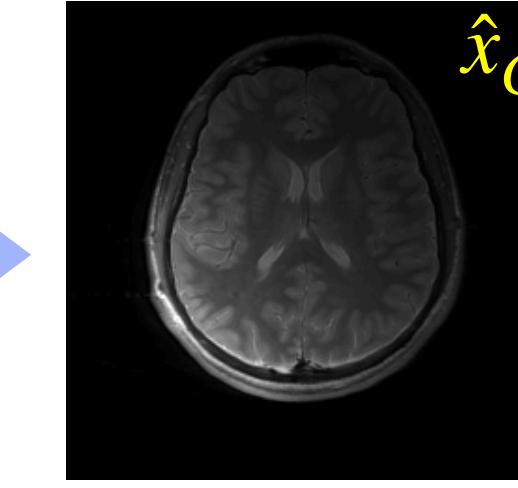
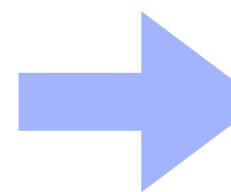
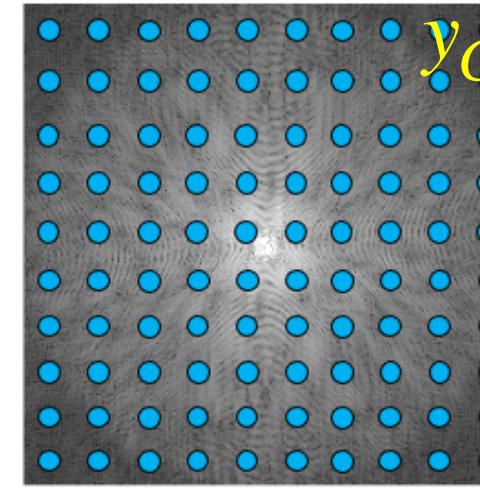
Reconstructed data

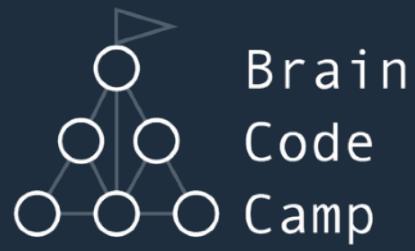


:

:

channel C





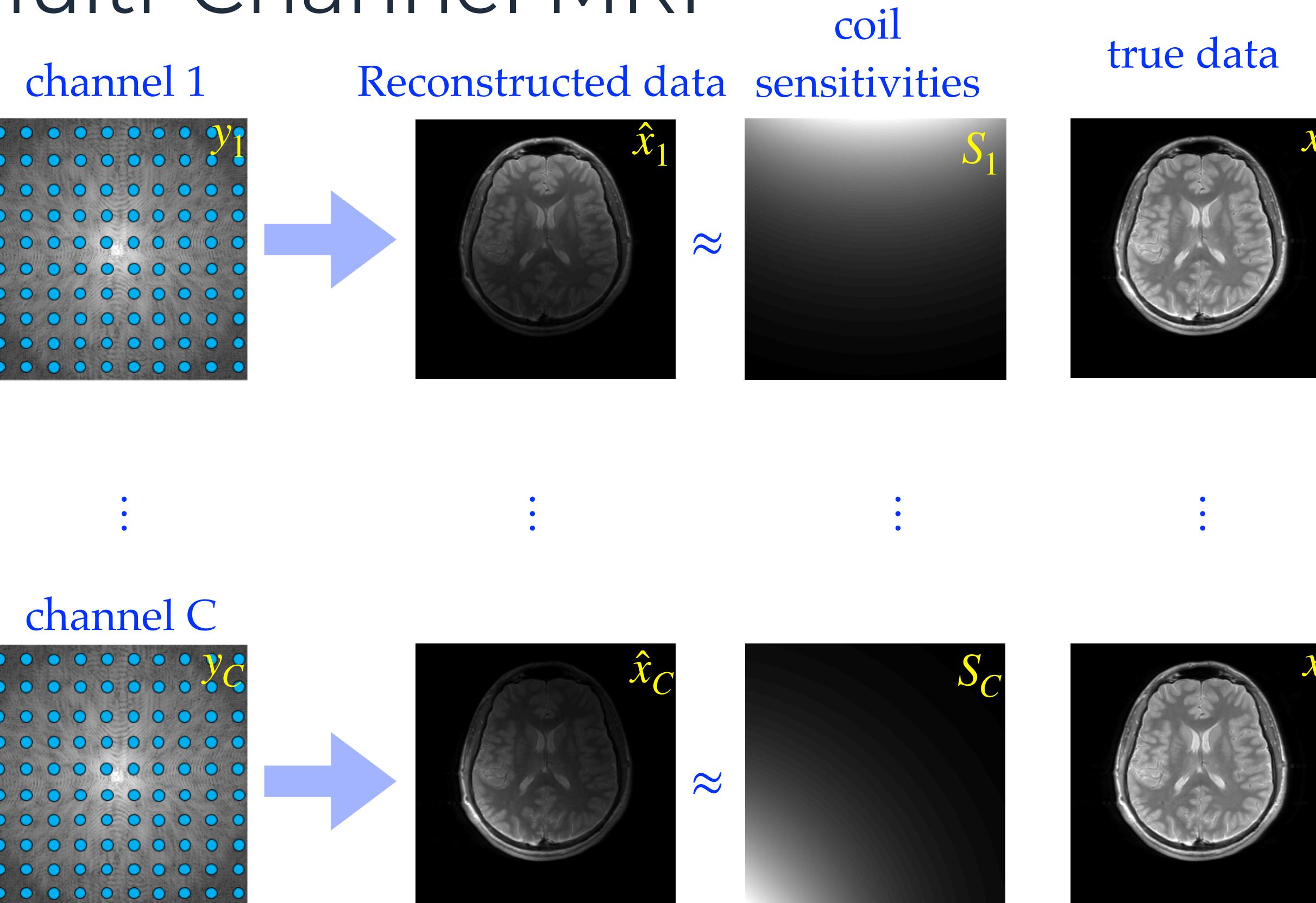
# Multi-Channel MRI



Keil et al. JMR (2013)

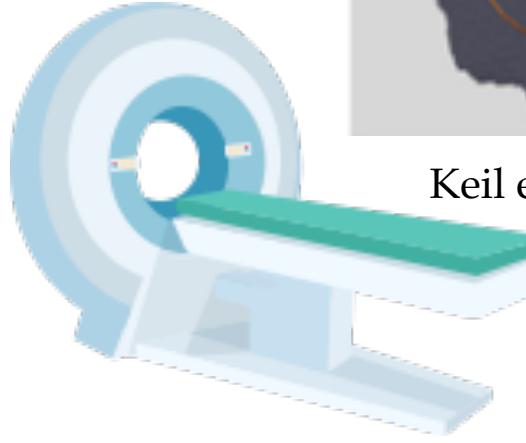
MR acquisition  
approximated using  
a linear operator

$$A \rightarrow AS_c$$



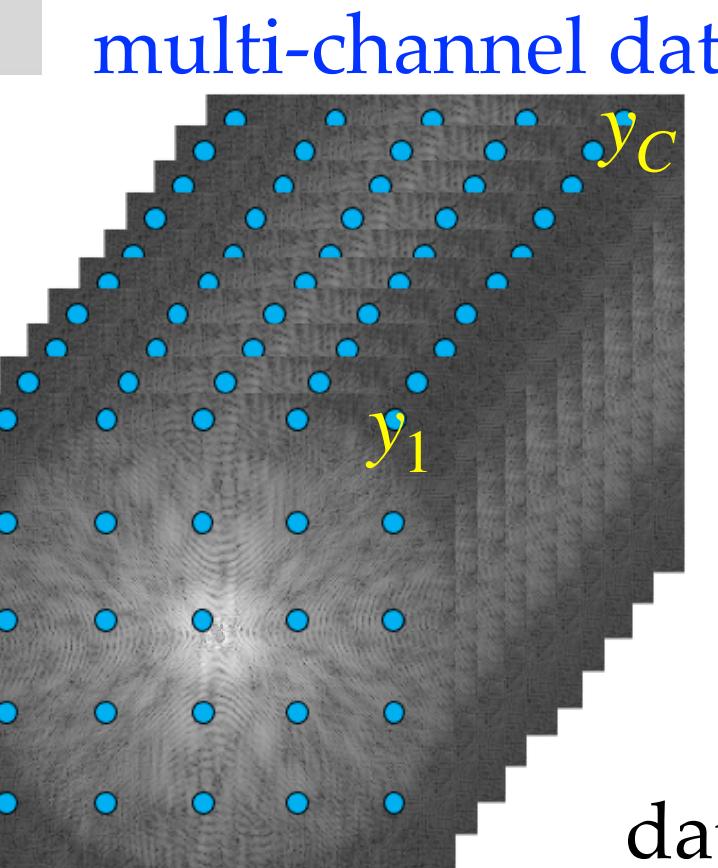
# Multi-Channel MRI

single-channel data



MR acquisition  
approximated using  
a linear operator

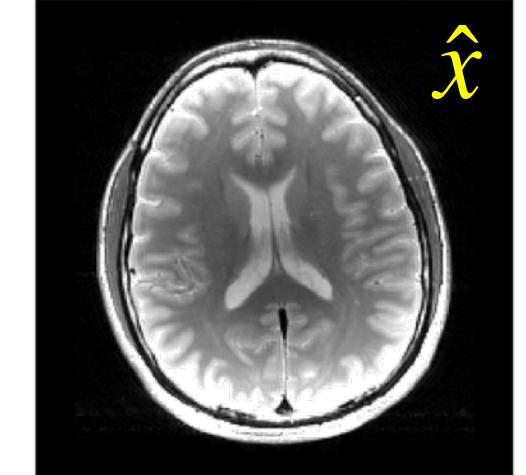
$$A \rightarrow AS_c$$



data dimension: 512 x 512 x 64 x 64 x T  
 row col slice channels time points

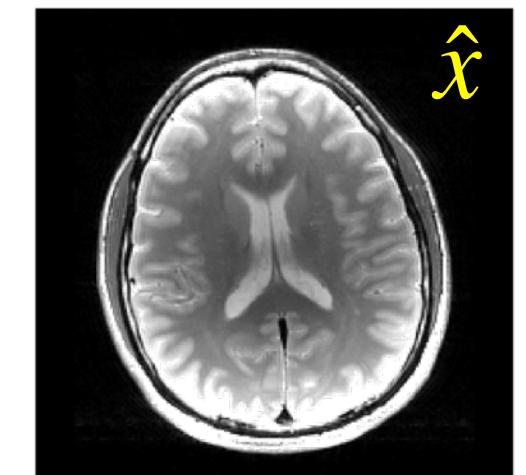
$$\min_x \frac{1}{2} \|Ax - y\|_2^2 + \lambda R(x)$$

single-channel reconstruction

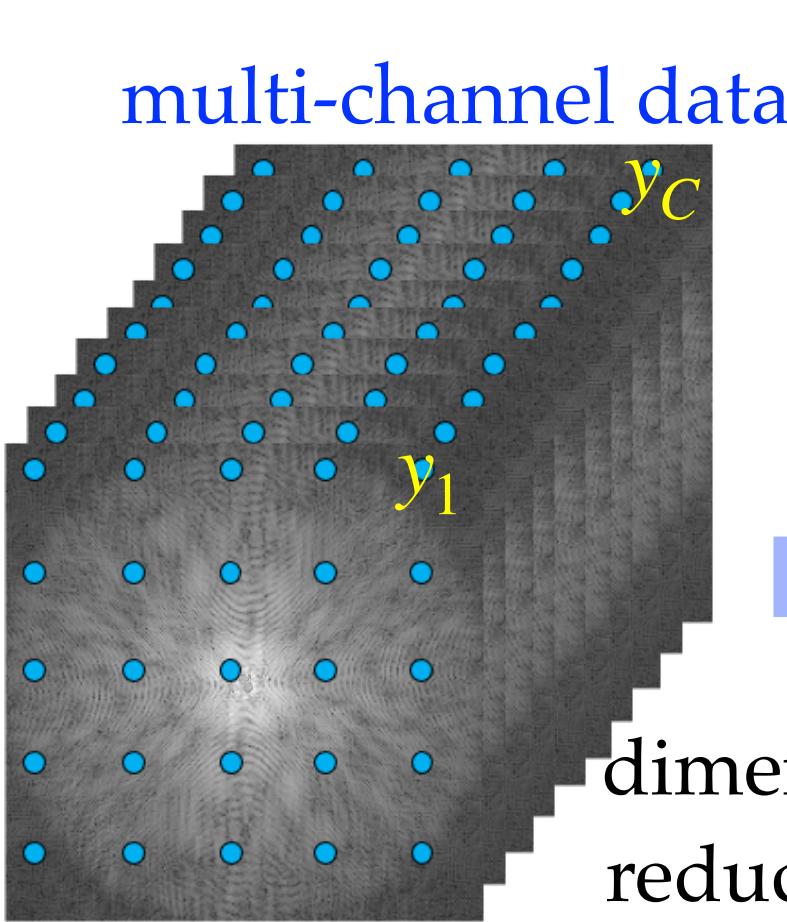


$$\min_x \frac{1}{2} \sum_c \|AS_c x - y_c\|_2^2 + \lambda \tilde{R}(x)$$

multi-channel reconstruction

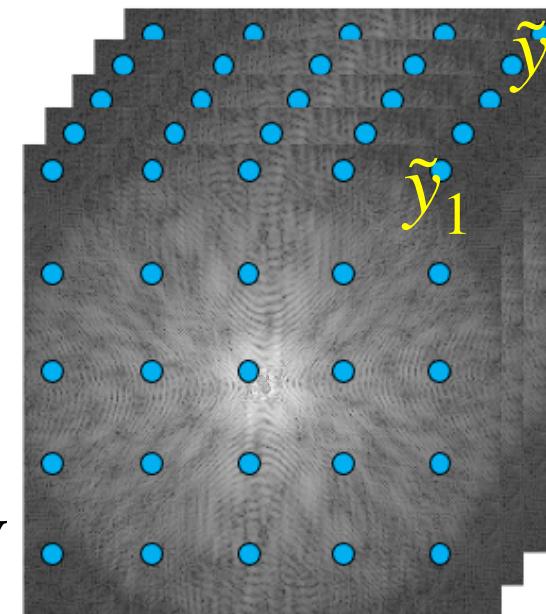


# Multi-Channel MRI with Virtual Coils



dimensionality reduction such as PCA

k-space data with fewer channels

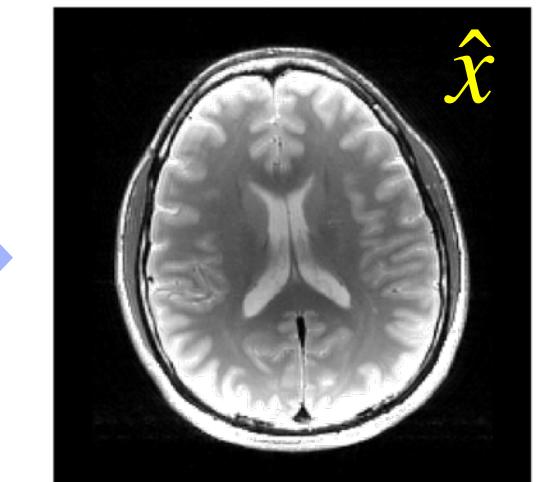


64 physical channels  
 $\rightarrow$  10 virtual channels

$$\min_x \frac{1}{2} \sum_c \|AS_c x - y_c\|_2^2 + \lambda \tilde{R}(x)$$

$$\min_x \frac{1}{2} \sum_c \|A\tilde{S}_c x - \tilde{y}_c\|_2^2 + \lambda \tilde{R}(x)$$

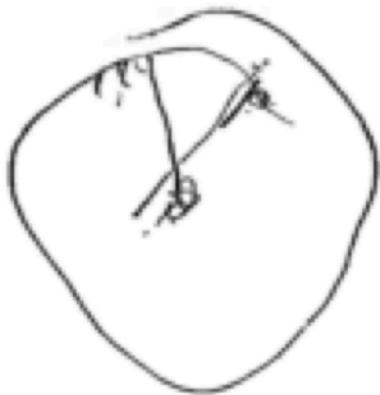
multi-channel reconstruction



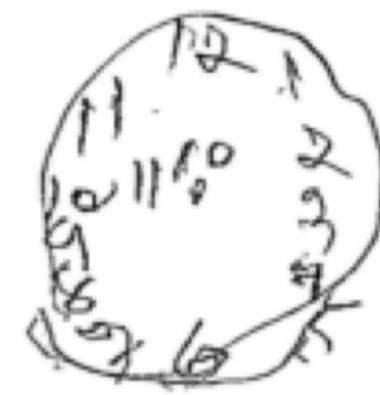
**Reduced computational resources**

# Clock Drawing Test

Score 0  
(n=13)



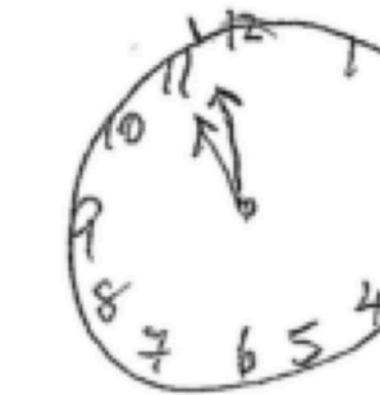
Score 1  
(n=20)



Score 2  
(n=53)



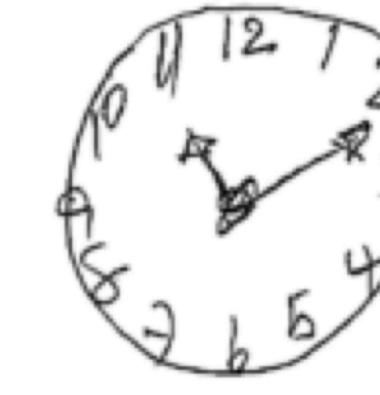
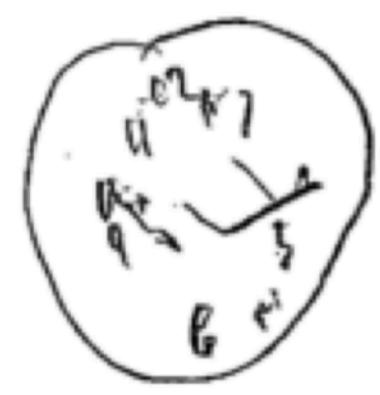
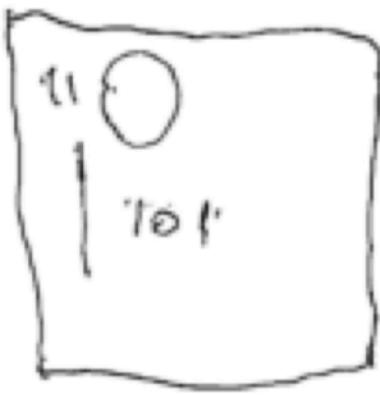
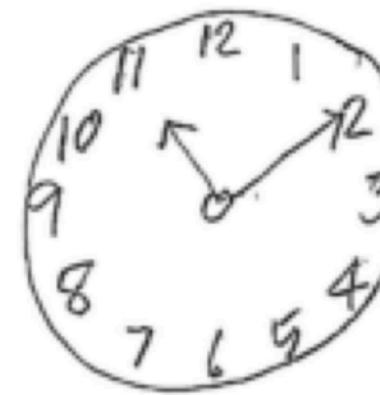
Score 3  
(n=352)



Score 4  
(n=1047)

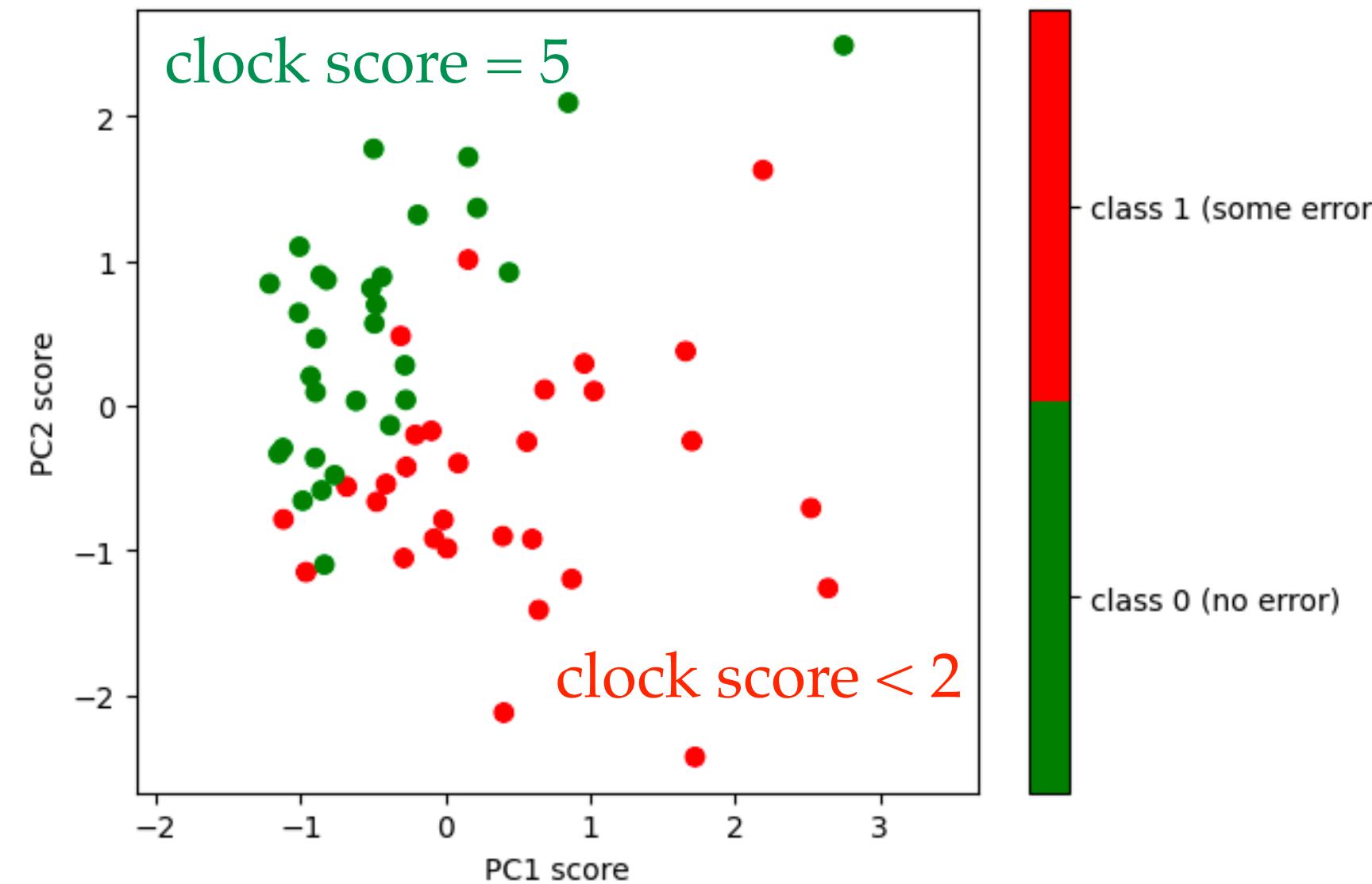


Score 5  
(n=1623)



# Clock Drawing Test

PCA



Contrastive learning + deep learning  
→ UMAP



Ruengchajatuporn, Natthanan, et al. "An explainable self-attention deep neural network for detecting mild cognitive impairment using multi-input digital drawing tasks." *Alzheimer's Research & Therapy* 14.1 (2022): 1-11.

Raksasat, Raksit, et al. "Attentive Pairwise Interaction Network for AI-assisted Clock Drawing Test Assessment of Early Visuospatial Deficits." Available at SSRN 4327538.