

# PSY9511: Seminar 5

## Unsupervised learning

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Esten H. Leonardsen

24.10.24



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# Outline

1. Exercise 4
2. Overview of unsupervised learning
3. Clustering
  - K-means
  - Hierarchical
4. Dimensionality reduction
  - Principal component analysis (PCA)
  - Independent component analysis (ICA)
  - Partial least squares (PLS)



# Exercise 4

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# Exercise 4: Stratification

<http://localhost:8888/notebooks/notebooks%2FStratification.ipynb>



# Exercise 4: Solution

<http://localhost:8888/notebooks/notebooks/Solution%204.ipynb>



# Unsupervised learning

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# Unsupervised learning: Motivation

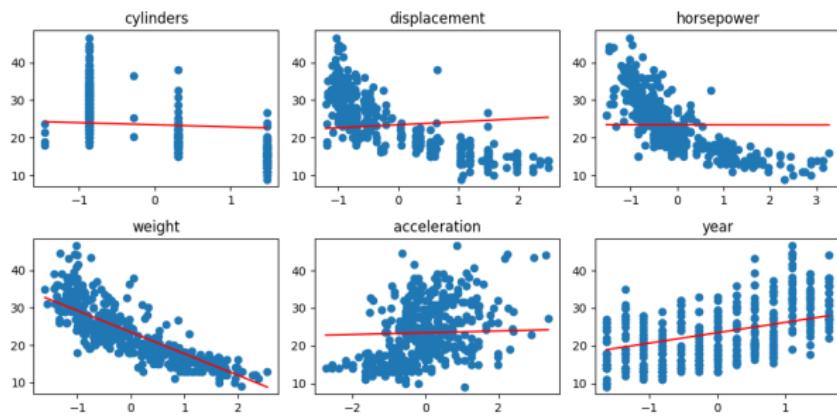
Supervised learning: Find  $\hat{y} = f(X)$



# Unsupervised learning: Motivation

## Supervised learning: Find $\hat{y} = f(X)$

- Descriptive: Understand the relationship between  $X$  and  $y$



# Unsupervised learning: Motivation

## Supervised learning: Find $\hat{y} = f(X)$

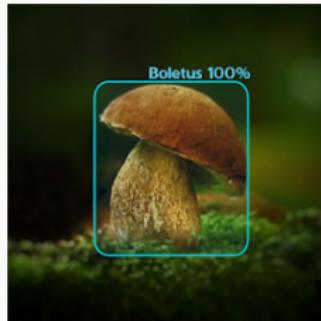
- Descriptive: Understand the relationship between  $X$  and  $y$
- Predictive: Predict  $y$  given new  $X$ .



# Unsupervised learning: Motivation

## Supervised learning: Find $\hat{y} = f(X)$

- Descriptive: Understand the relationship between  $X$  and  $y$
- Predictive: Predict  $y$  given new  $X$ .
  - Because the predictions are useful



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## Unsupervised learning: Are there interesting patterns in $X$ ?



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## Unsupervised learning: Are there interesting patterns in $X$ ?

- Can we find subgroups or interesting axes of variability?



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## Supervised learning: Find $\hat{y} = f(X)$

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## Unsupervised learning: Are there interesting patterns in $X$ ?

- Can we find subgroups or interesting axes of variability?
- Exploratory analyses
- Visualization



# Clustering

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# Clustering: Background

Are there some (naturally occurring) subgroups in our dataset?



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Are there some (naturally occurring) subgroups in our dataset?

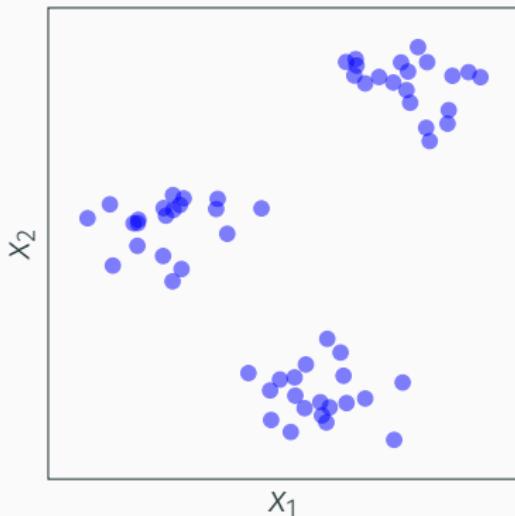
$x_1$	$x_2$
0.20	-0.26
0.15	-0.33
0.03	0.07
-0.07	-0.01
-0.06	0.00
0.28	-0.24
0.21	-0.35
0.20	-0.32
0.30	0.25
0.00	-0.12



# Clustering: Background

Are there some (naturally occurring) subgroups in our dataset?

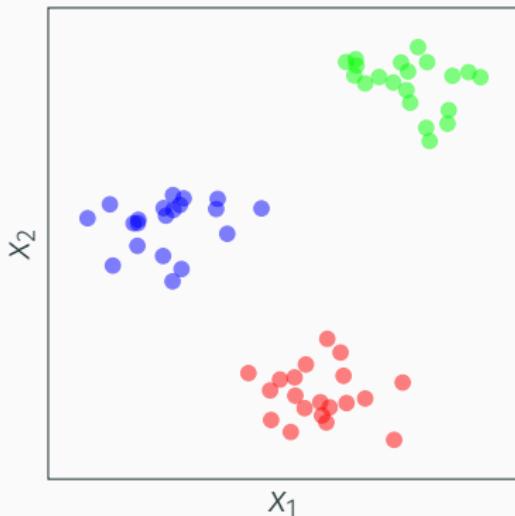
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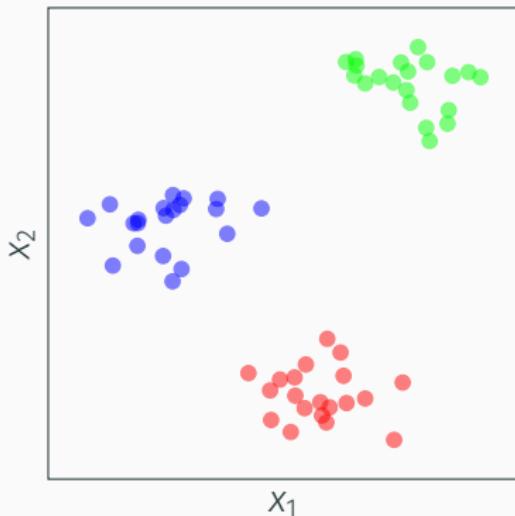
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-0.06	0.00
0.28	-0.24
0.21	-0.35
0.20	-0.32
0.30	0.25
0.00	-0.12



# K-means clustering: Definition

K-means clustering: Find  $k$  clusters in the data to minimize the *within-cluster variance*:

$$\underset{C_1, \dots, C_k}{\text{minimize}} \left( \sum_{k=1}^K \frac{1}{|C_k|} \sum_{i, i' \in C_k} \sum_{j=1}^p (x_{ij} - x_{i'j})^2 \right)$$



# K-means clustering: Definition

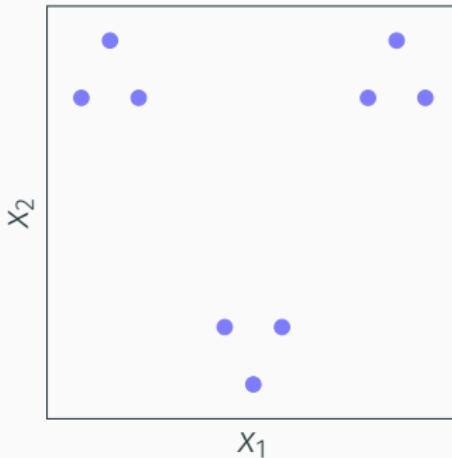
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# K-means clustering: Definition

$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25

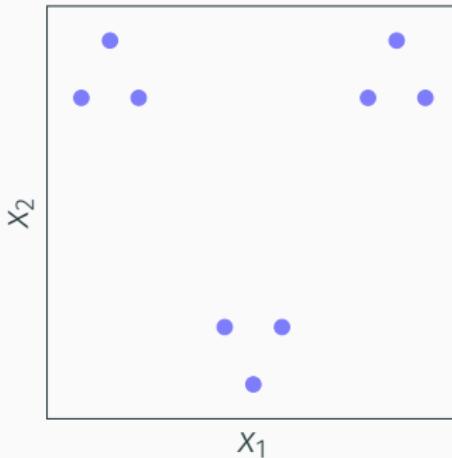


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# K-means clustering: Definition

$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25



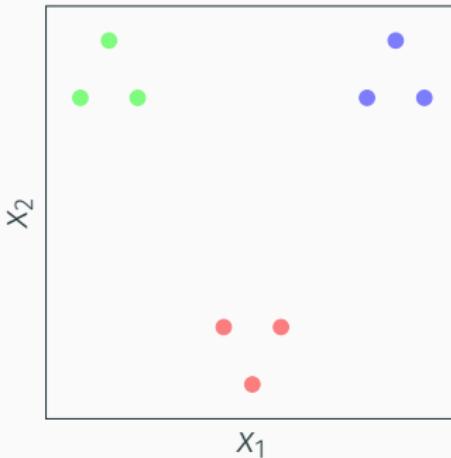
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# K-means clustering: Definition

$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25

$C_1$   
 $C_1$   
 $C_1$   
 $C_2$   
 $C_2$   
 $C_2$   
 $C_3$   
 $C_3$   
 $C_3$



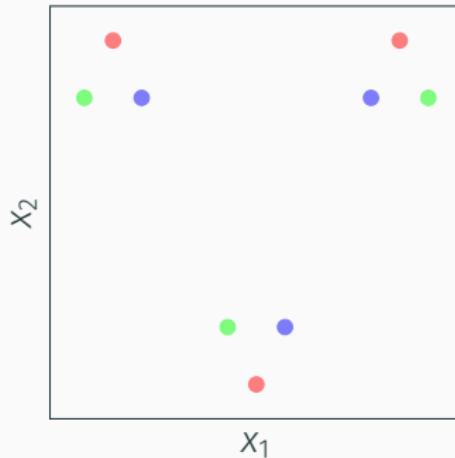
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# K-means clustering: Definition

$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25

$C_1$   
 $C_2$   
 $C_3$   
 $C_1$   
 $C_2$   
 $C_3$   
 $C_1$   
 $C_2$   
 $C_3$



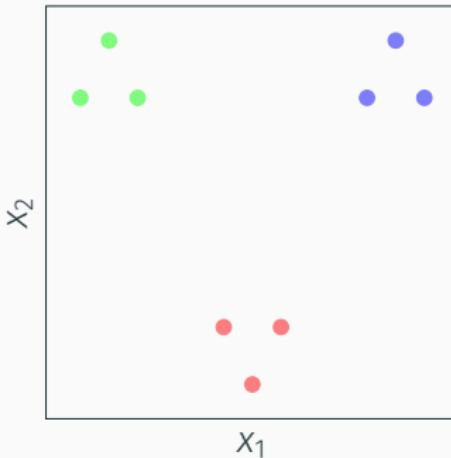
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# K-means clustering: Definition

$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25

$C_1$   
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 $C_2$   
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 $C_3$   
 $C_3$   
 $C_3$



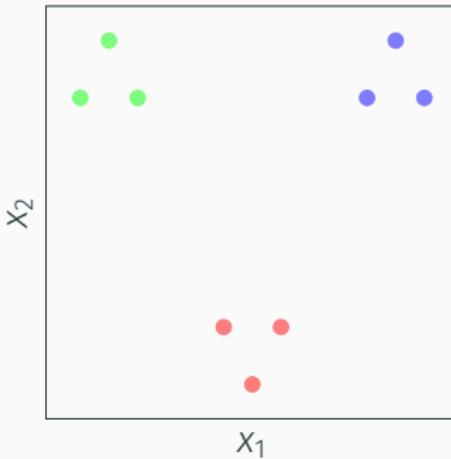
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1	1
1.5	1
1.25	1.25
-1	1
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-1.25	1.25
0.25	0
-0.25	0
0	-0.25

$C_1$   
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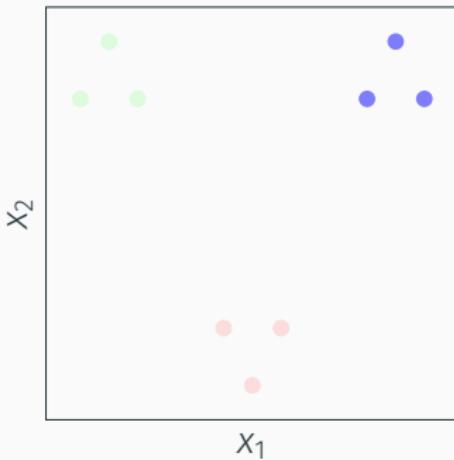


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$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
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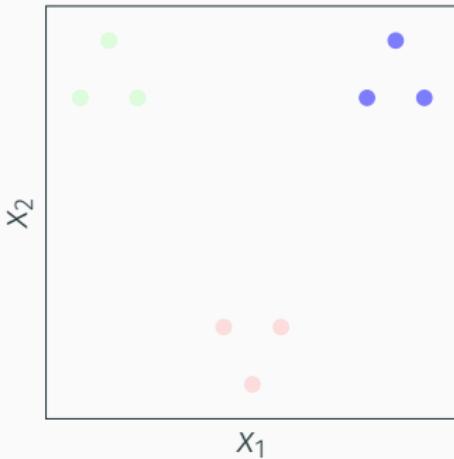
 $C_1$  $C_1$  $C_1$  $C_2$  $C_2$  $C_2$  $C_3$  $C_3$  $C_3$  $C_3$ 

$$\underset{C_1, \dots, C_k}{\text{minimize}} \left( \sum_{k=1}^K \frac{1}{|C_k|} \sum_{i, i' \in C_k} \sqrt{\sum_{j=1}^p (x_{ij} - x_{i'j})^2} \right)$$



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1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25

 $C_1$  $C_1$  $C_1$  $C_2$  $C_2$  $C_2$  $C_3$  $C_3$  $C_3$  $C_3$ 

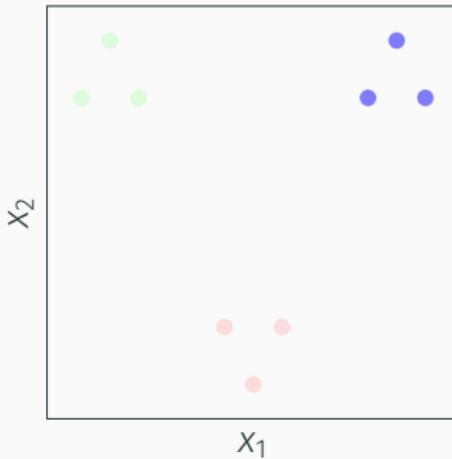
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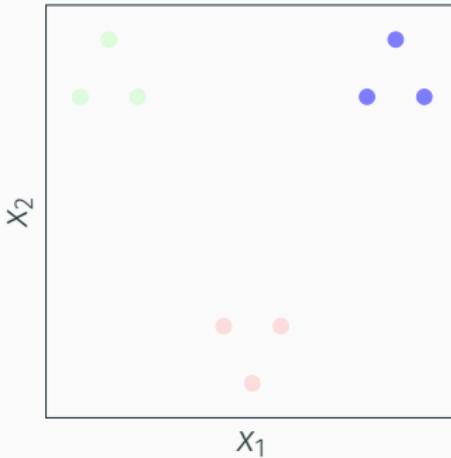
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# K-means clustering: Definition

	$x_1$	$x_2$
$\uparrow$ $i, i'$ $\downarrow$		
1	1	1
1.5	1	
1.25	1.25	
-1	1	
-1.5	1	
-1.25	1.25	
0.25	0	
-0.25	0	
0	-0.25	

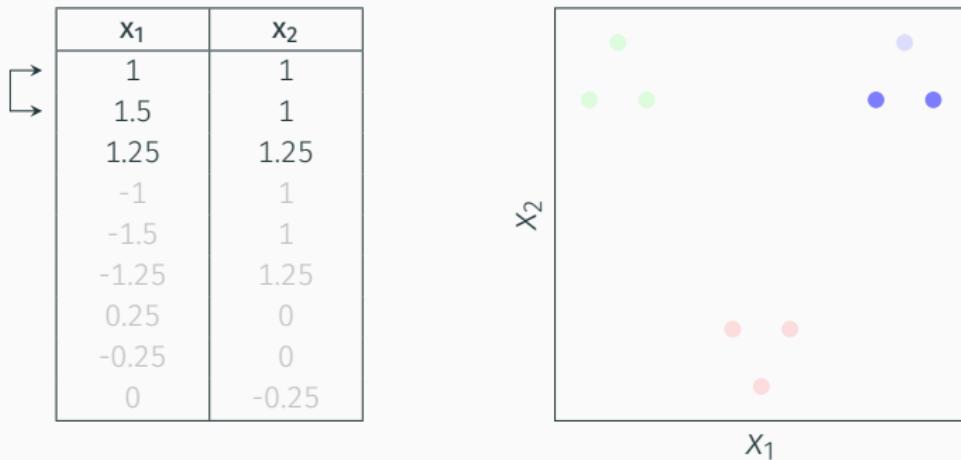
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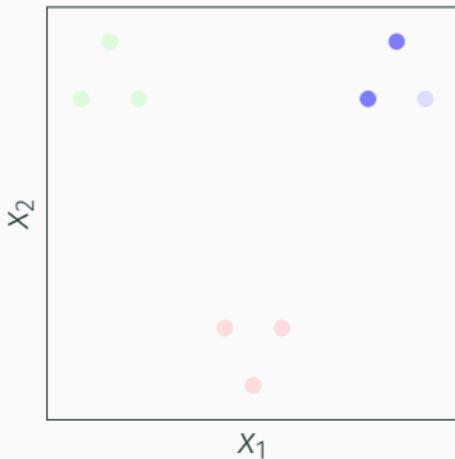
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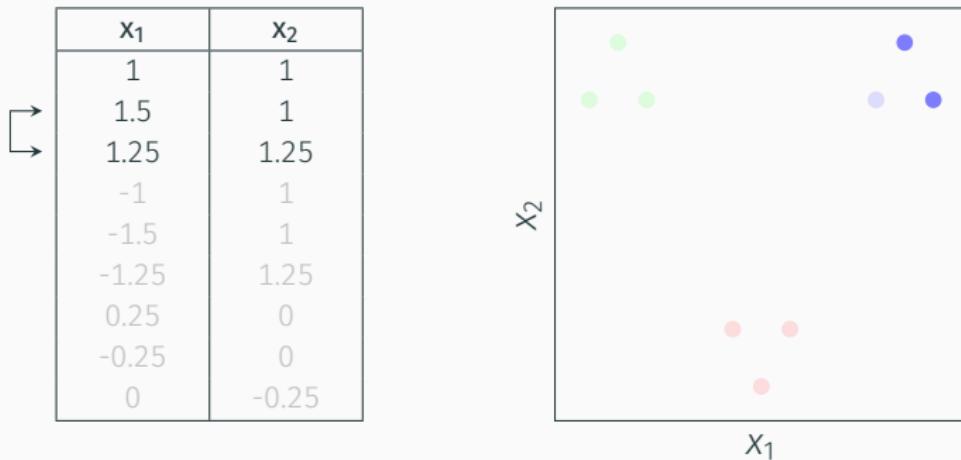
$x_1$	$x_2$
1	1
1.5	1
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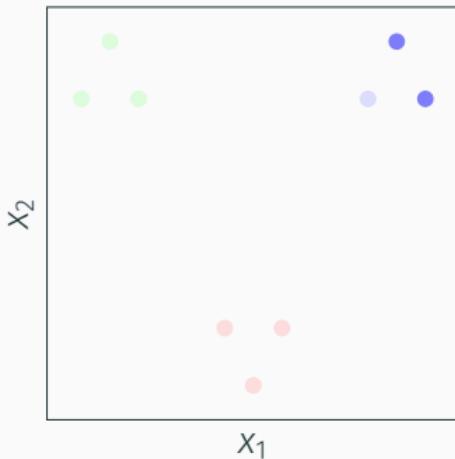


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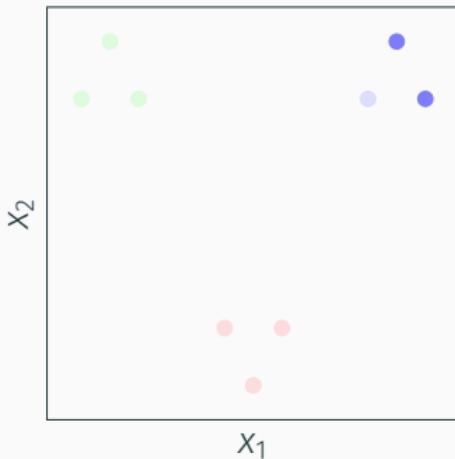


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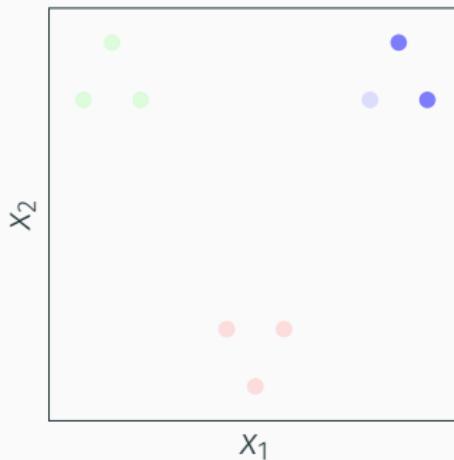
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↔ j →

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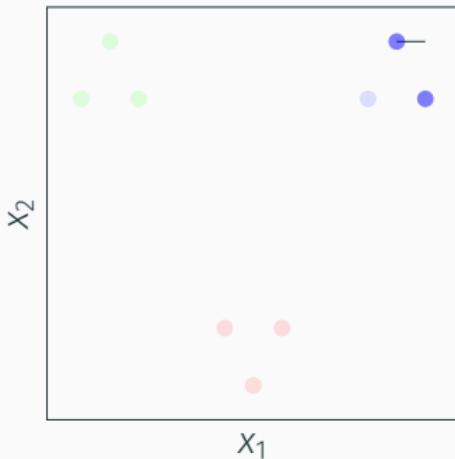


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0.25	0
-0.25	0
0	-0.25

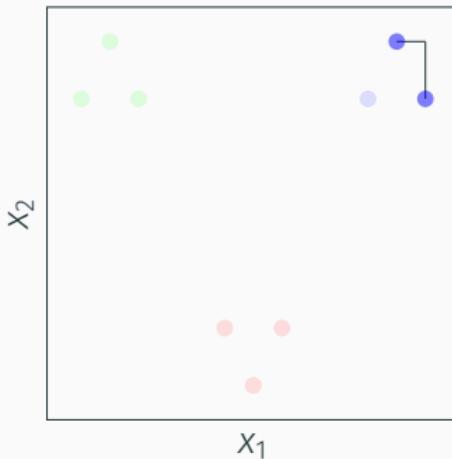


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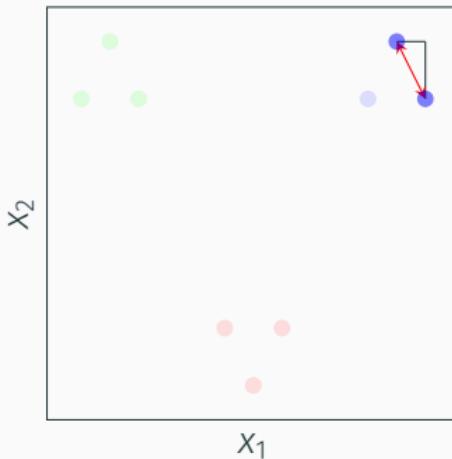


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# K-means clustering: Definition

We want to find an assignment of clusters that for each cluster, for each pair of points within the cluster, minimizes the distance between them

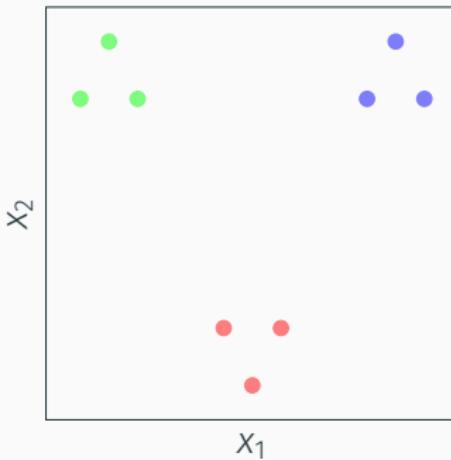
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# K-means clustering: Definition

$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25

$C_1$   
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 $C_2$   
 $C_2$   
 $C_2$   
 $C_3$   
 $C_3$   
 $C_3$



# K-means clustering: Definition

$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25

$C_1$

$C_2$

$C_3$

$C_1$

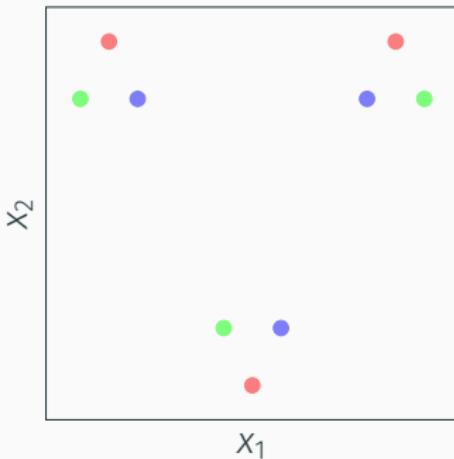
$C_2$

$C_3$

$C_1$

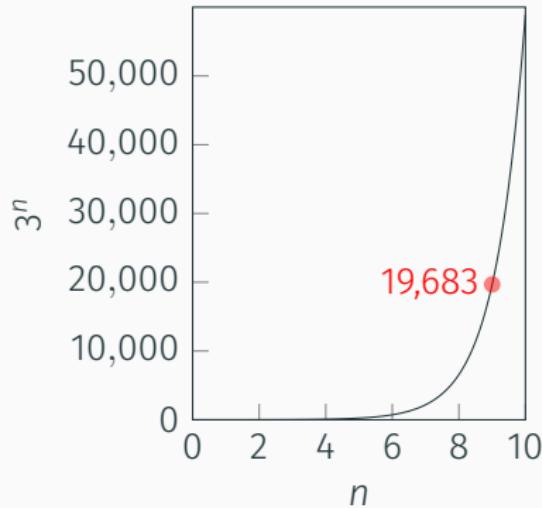
$C_2$

$C_3$



# K-means clustering: Definition

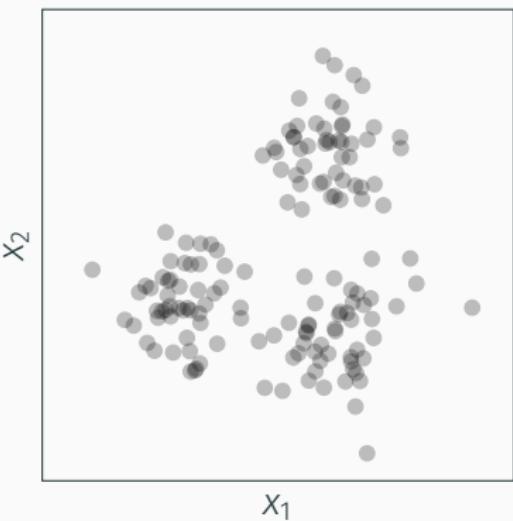
$x_1$	$x_2$
1	1
1.5	1
1.25	1.25
-1	1
-1.5	1
-1.25	1.25
0.25	0
-0.25	0
0	-0.25



Number of possible assignments:  $K^n$

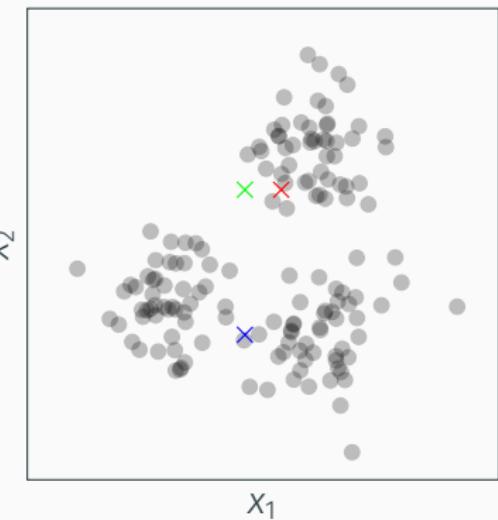


# K-means clustering: Algorithm



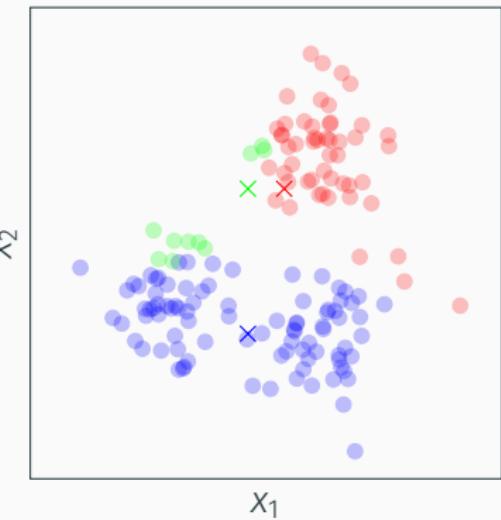
# K-means clustering: Algorithm

1. Initialize  $k$  random centroids



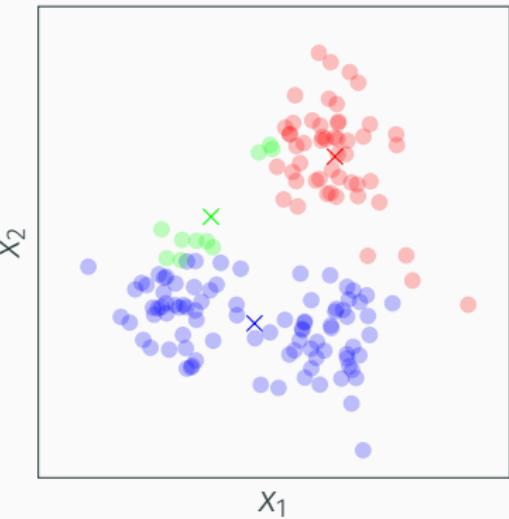
# K-means clustering: Algorithm

1. Initialize  $k$  random centroids
2. Iteratively:
  - 2.1 Assign each point to the nearest centroid



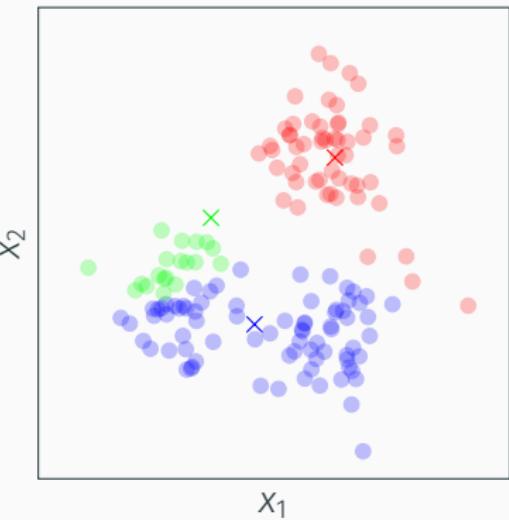
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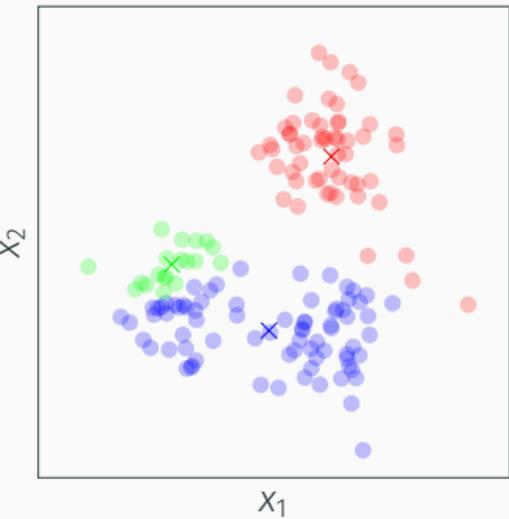
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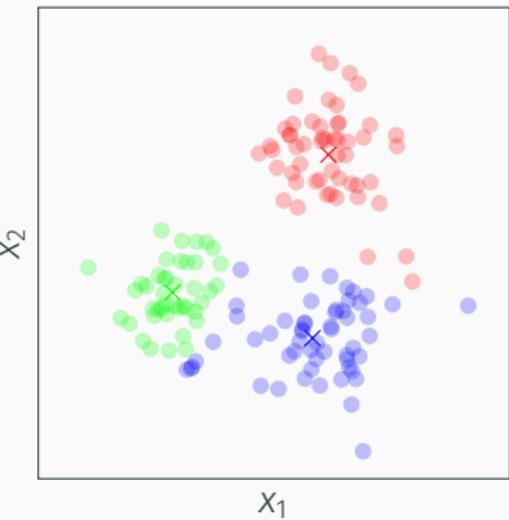
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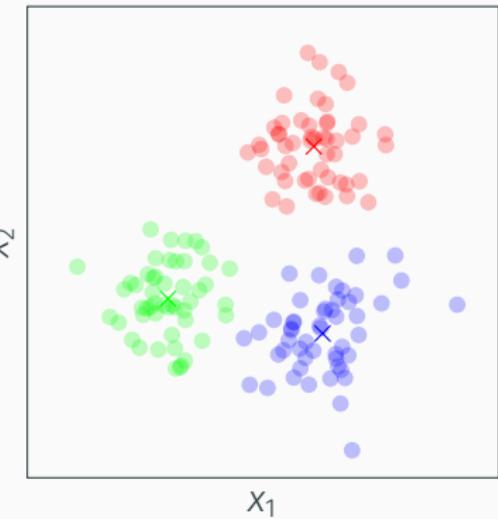
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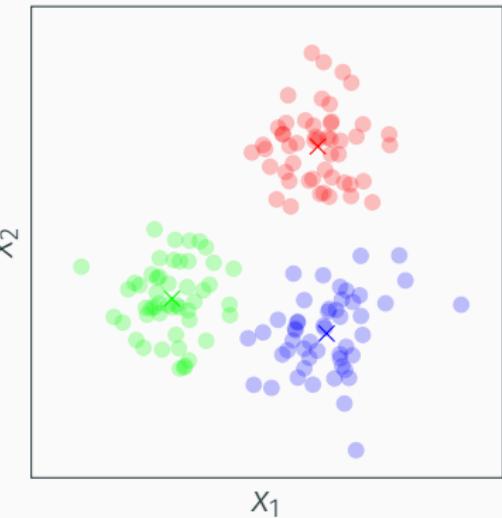
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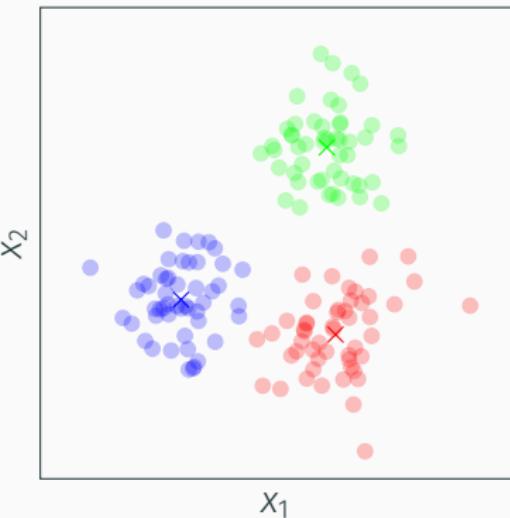
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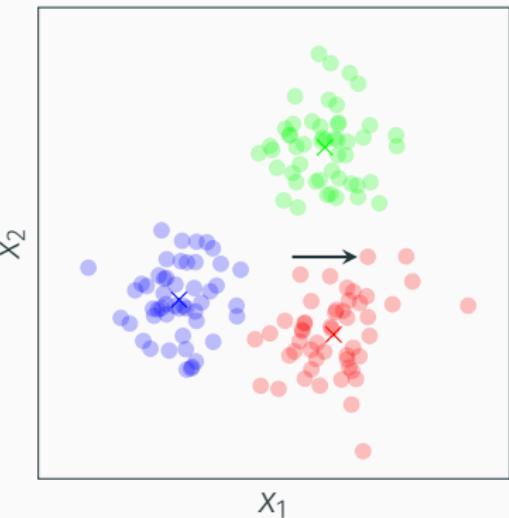
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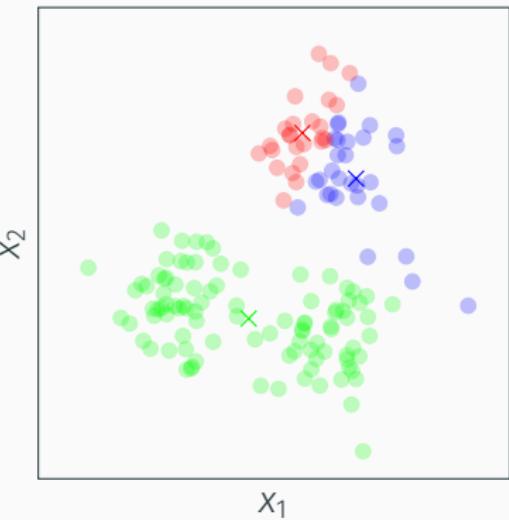
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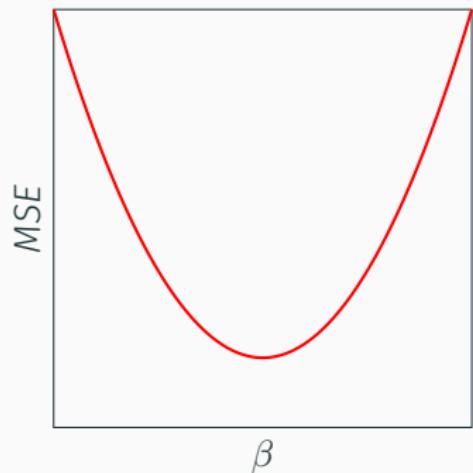


# K-means clustering: Algorithm

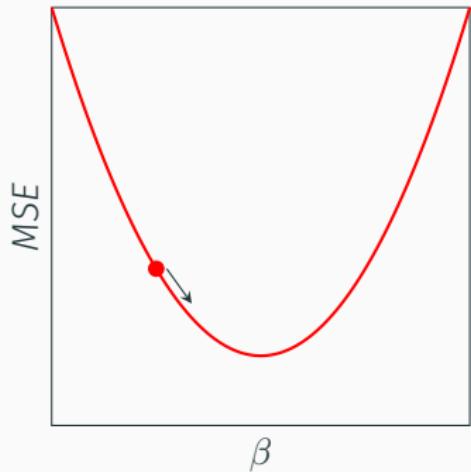
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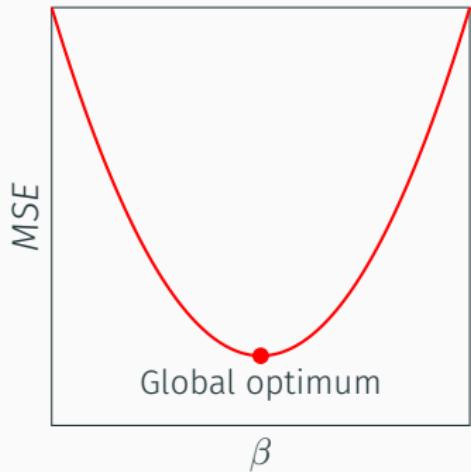
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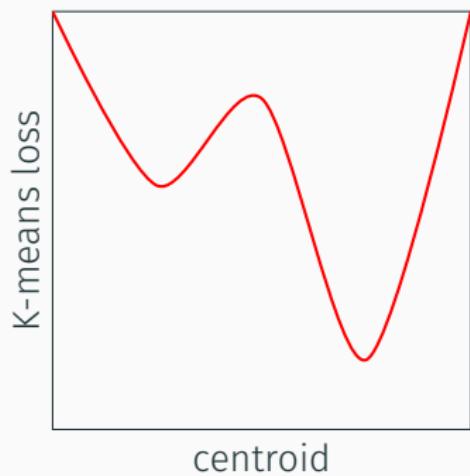
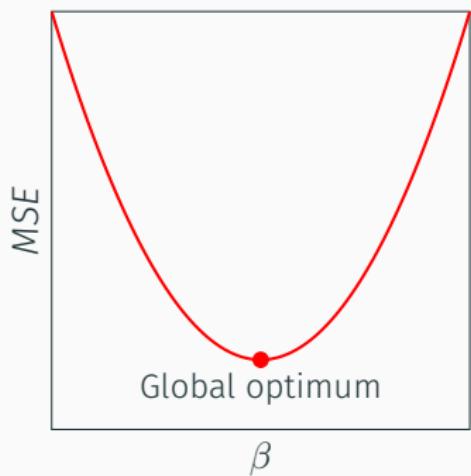
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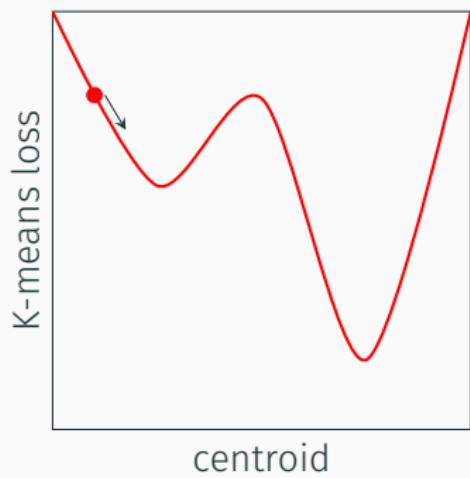
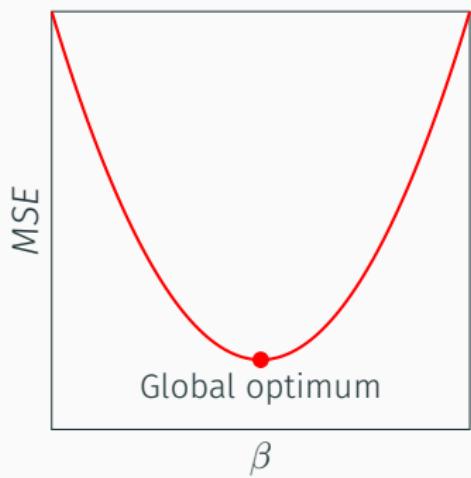
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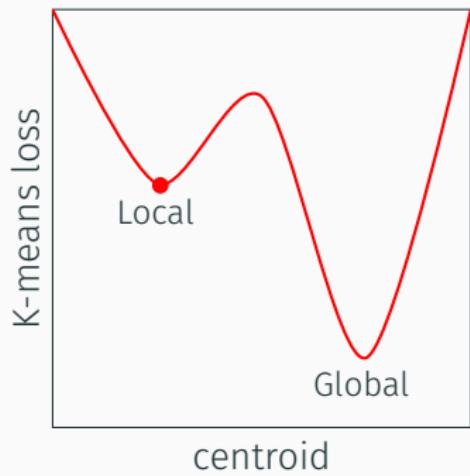
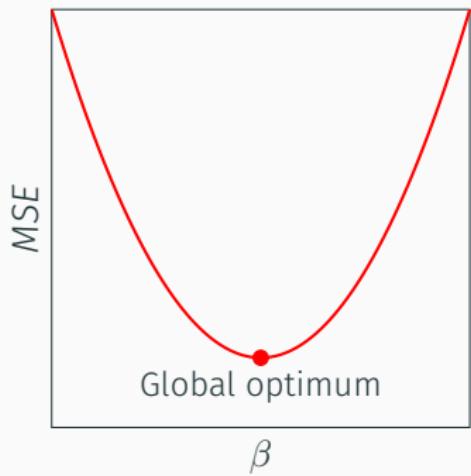
# K-means clustering: Algorithm



# K-means clustering: Algorithm

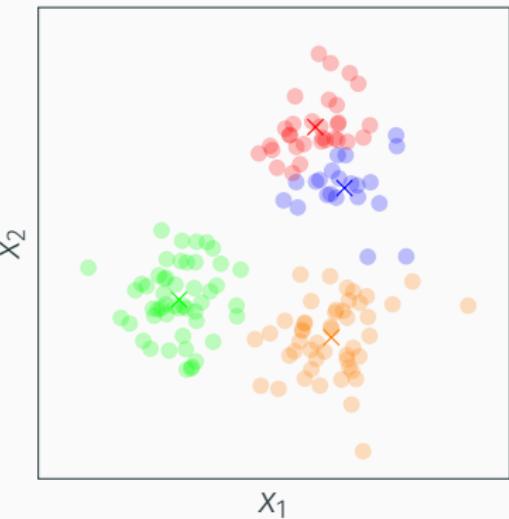


# K-means clustering: Algorithm



# K-means clustering: Algorithm

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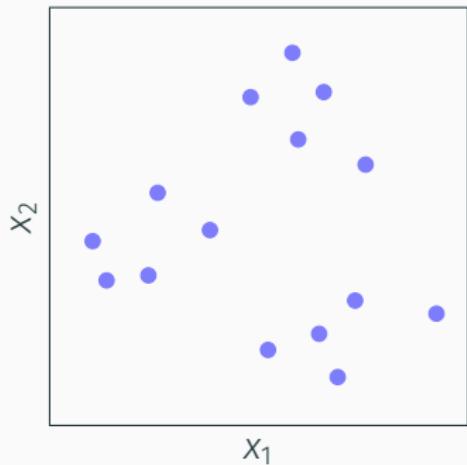
# K-means clustering: Summary

K-means clustering: Finds clusters by minimizing the within-cluster variance

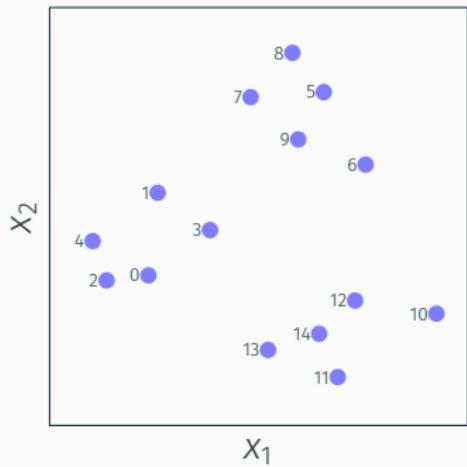
- + Intuitive way of finding clusters
- + Fast algorithm
- Dependent on knowing the number of clusters to use
- Dependent on the random initialization of centroids



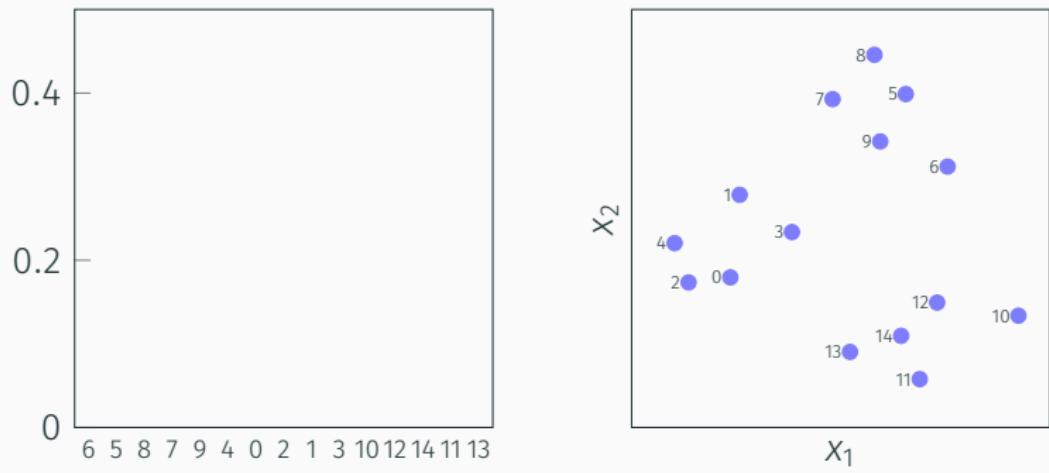
# Hierarchical clustering: Algorithm



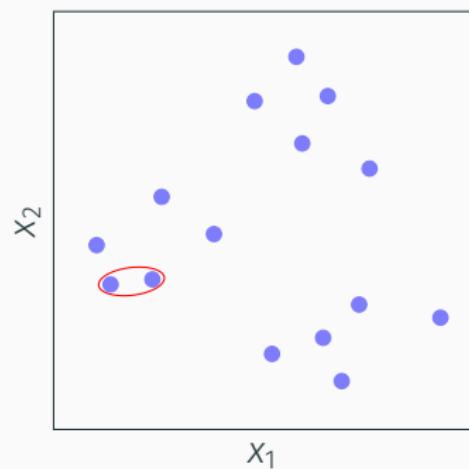
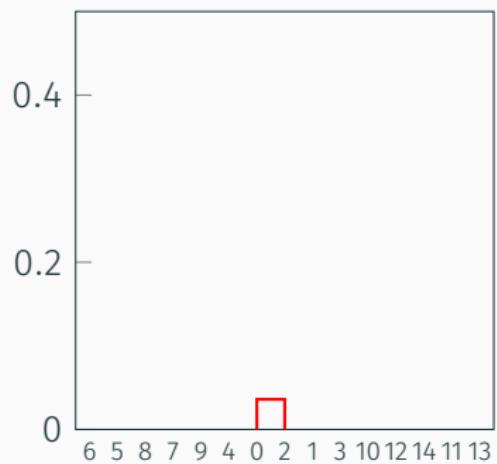
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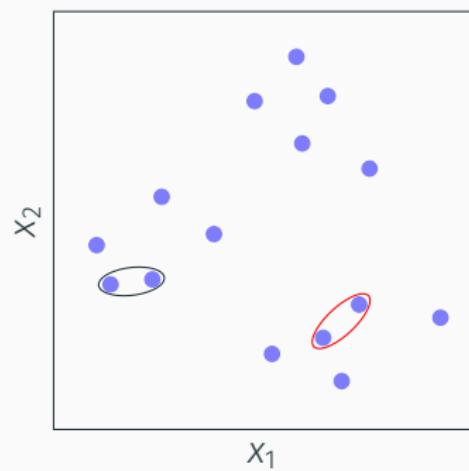
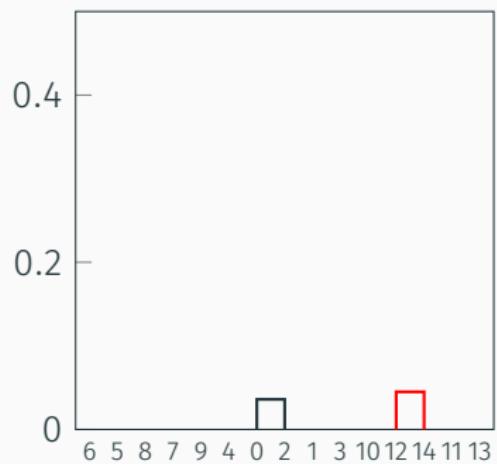
# Hierarchical clustering: Algorithm



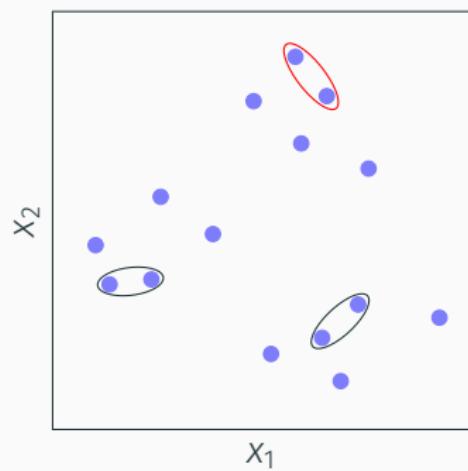
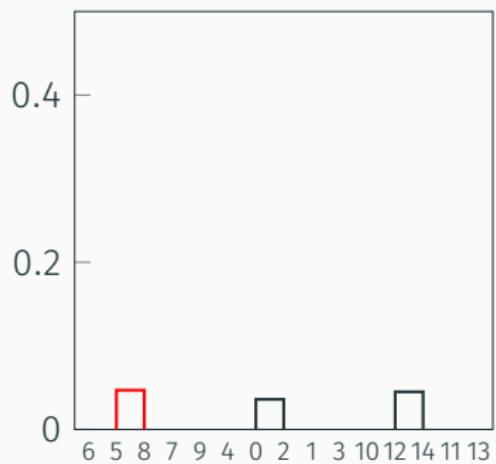
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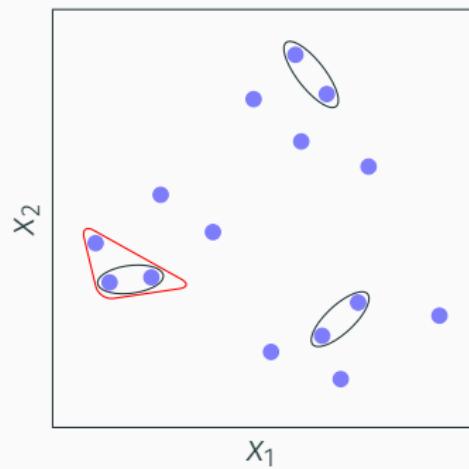
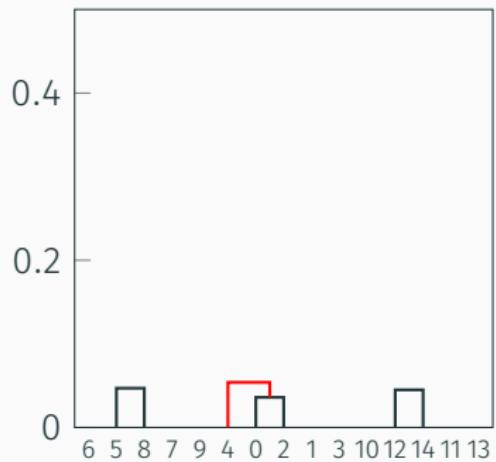
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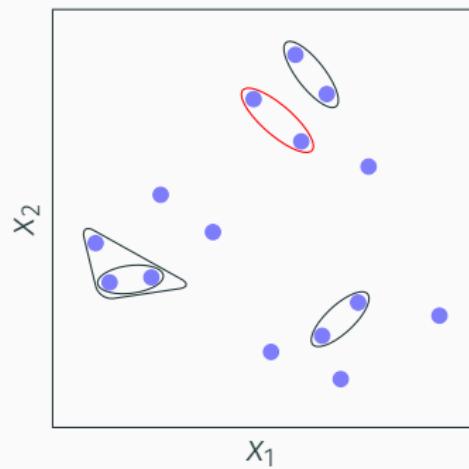
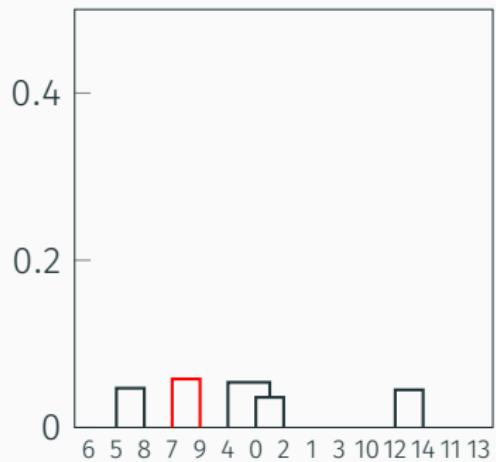
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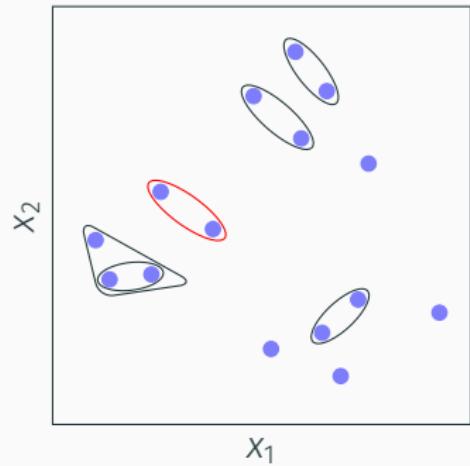
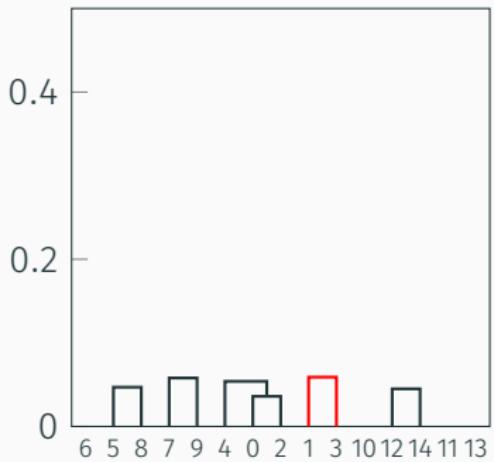
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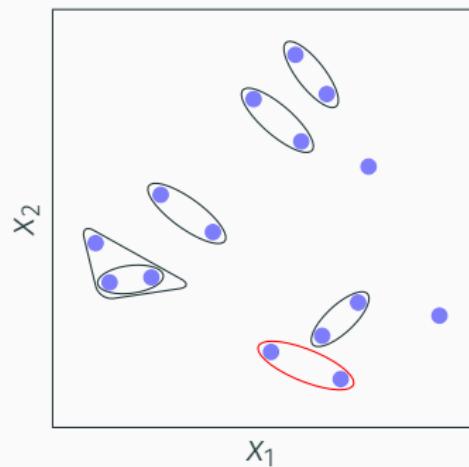
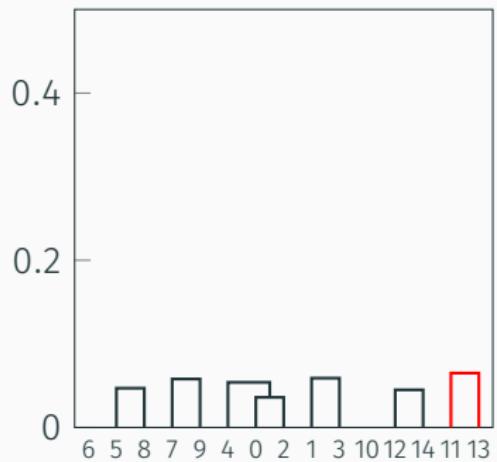
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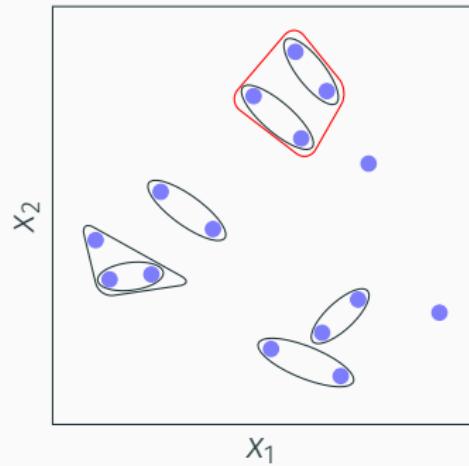
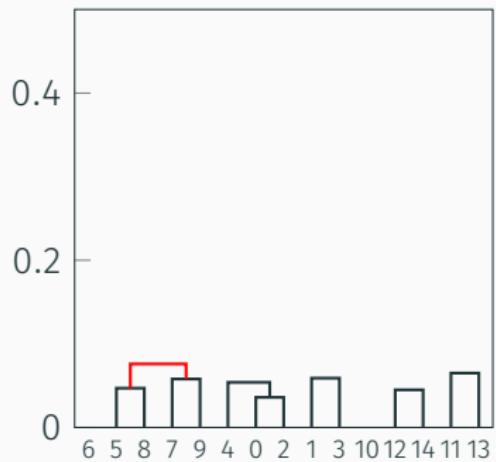
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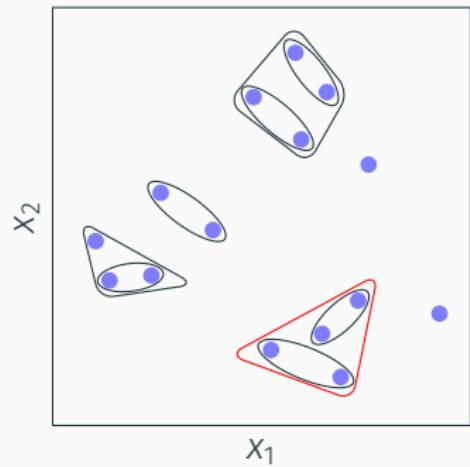
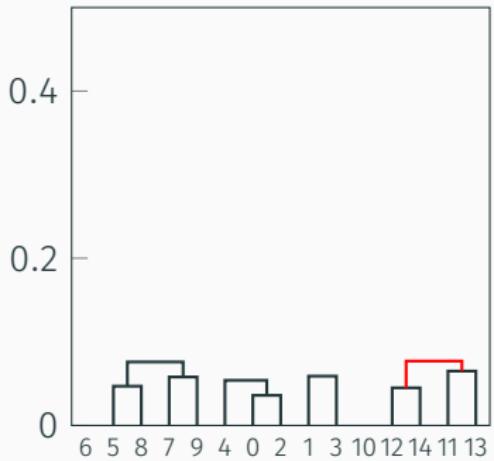
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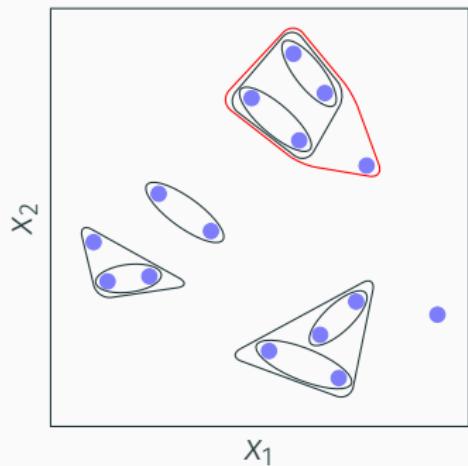
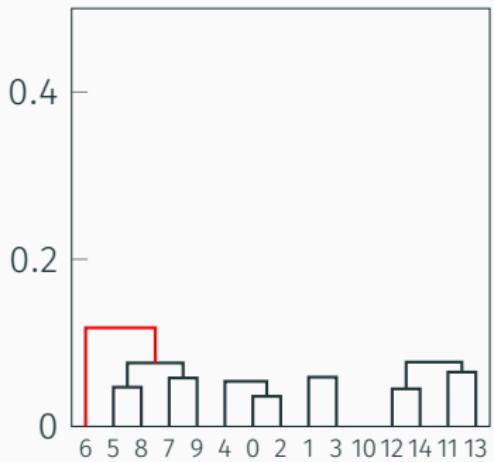
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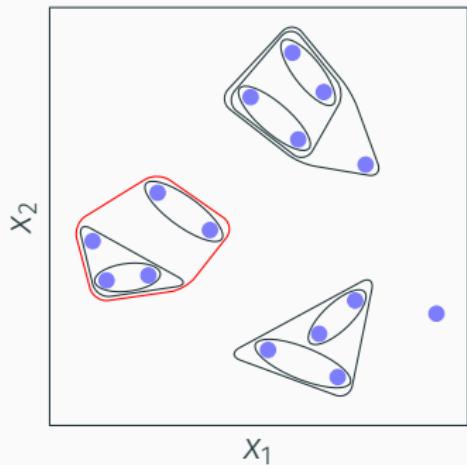
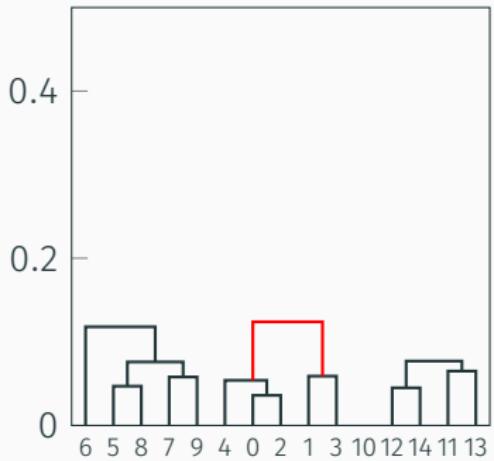
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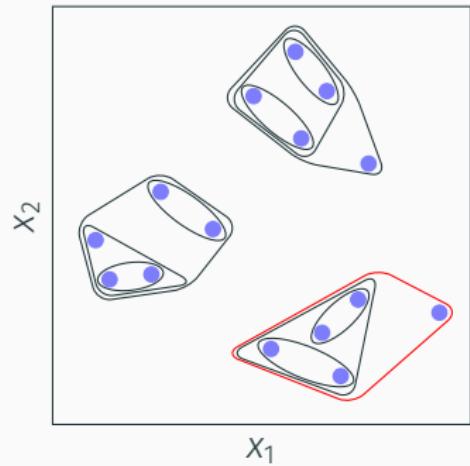
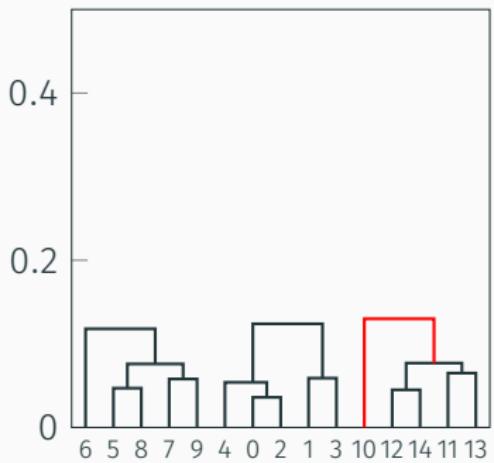
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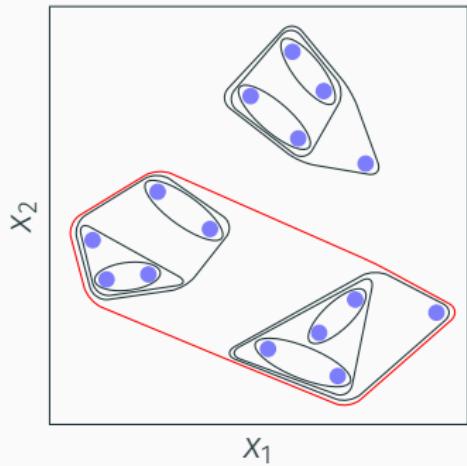
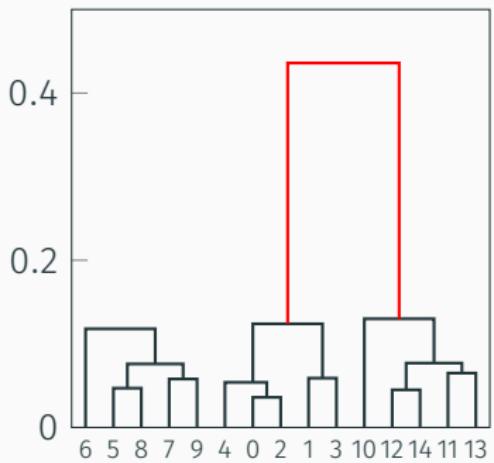
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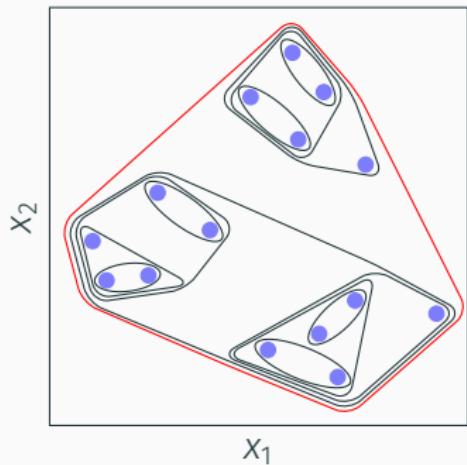
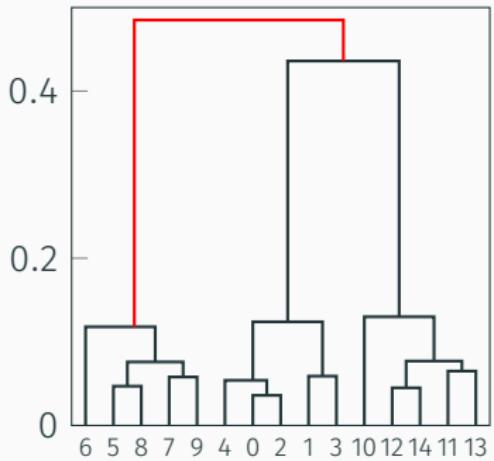
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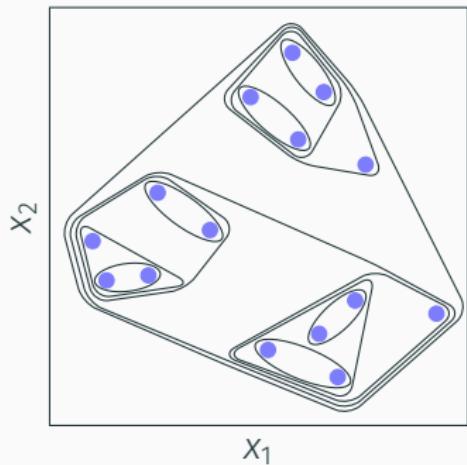
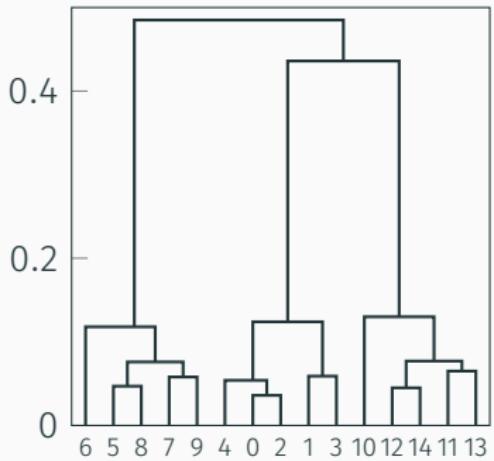
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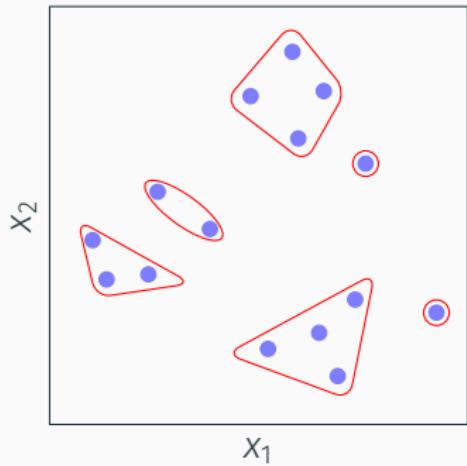
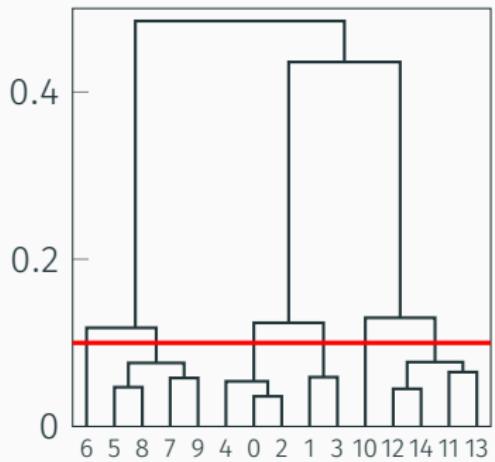
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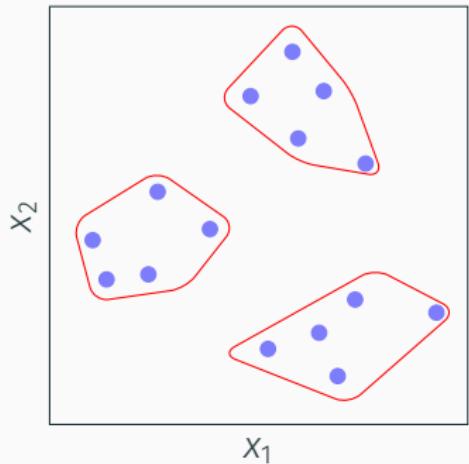
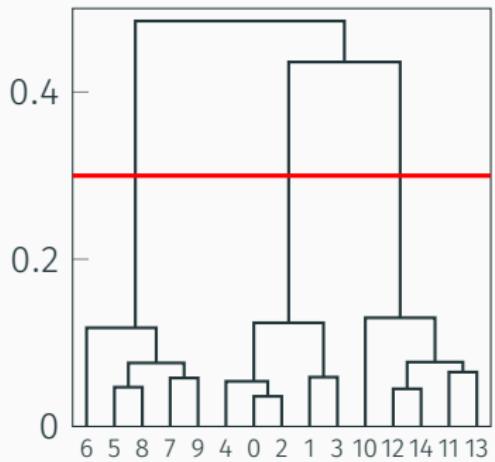
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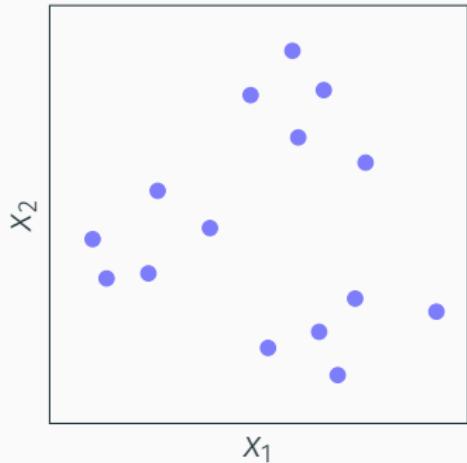
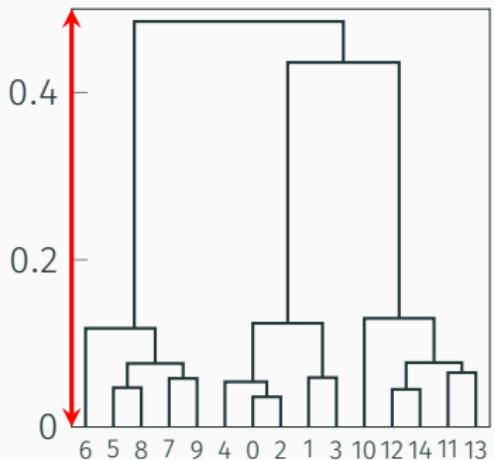
# Hierarchical clustering: Interpretation



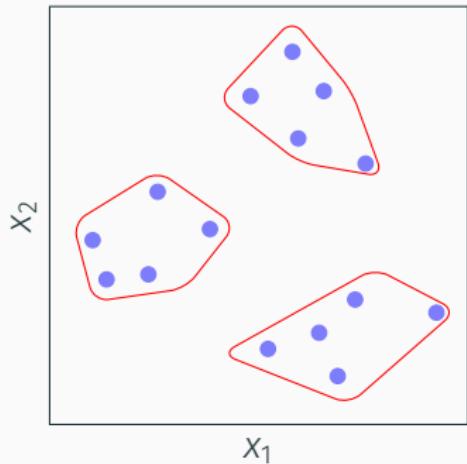
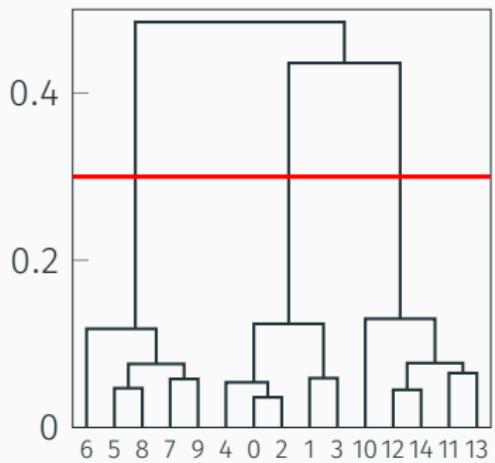
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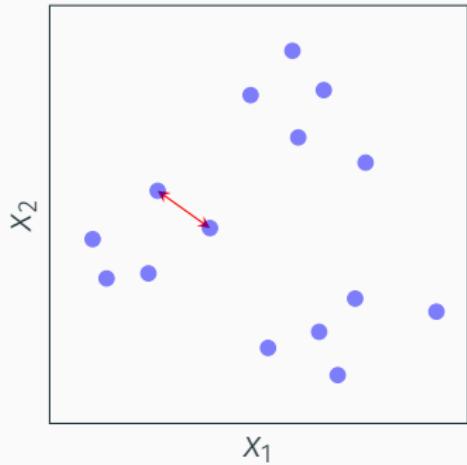
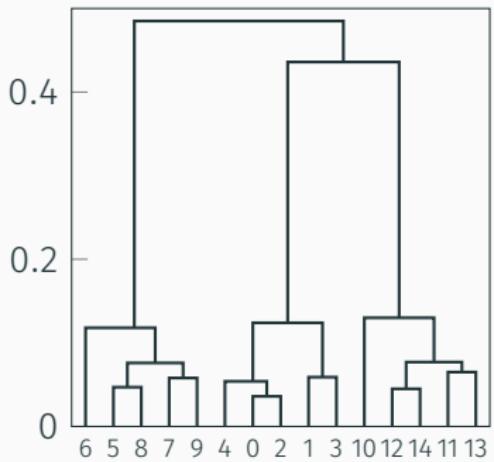
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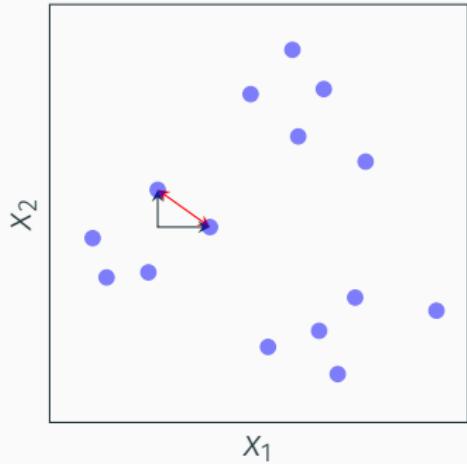
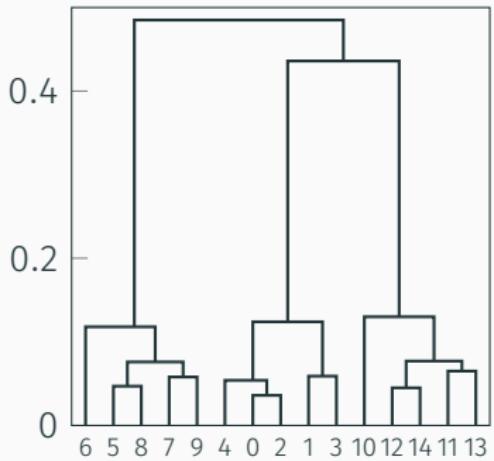
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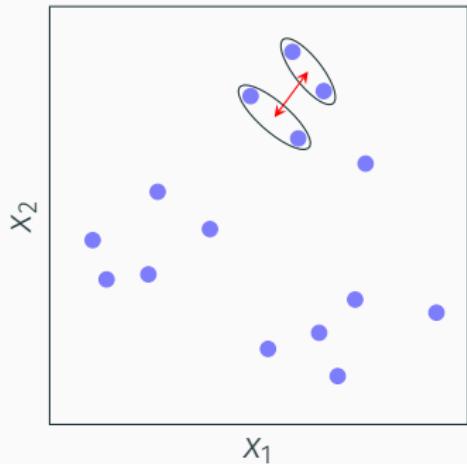
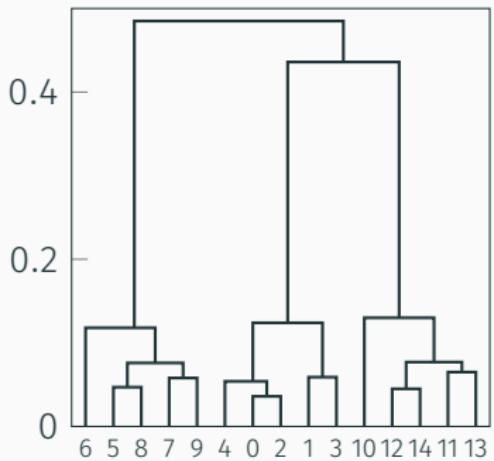
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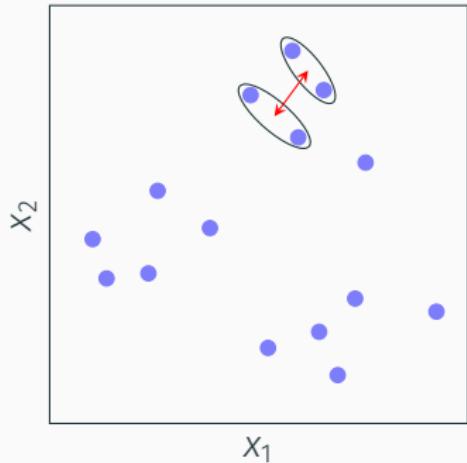
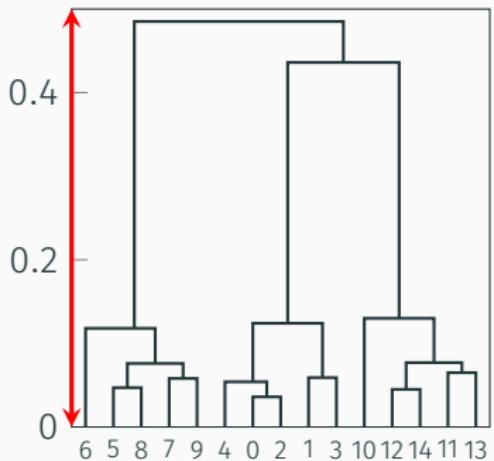
# Hierarchical clustering: Interpretation



# Hierarchical clustering: Interpretation



# Hierarchical clustering: Interpretation



# Hierarchical clustering: Interpretation

Complete	Maximal intercluster dissimilarity
Single	Minimal intercluster dissimilarity
Average	Mean intercluster dissimilarity
Centroid	Dissimilarity between the centroid for cluster A and the centroid for cluster B.



# Hierarchical clustering: Interpretation

Agglomerative clustering: Iteratively merge clusters to form a hierarchy of cluster assignments.

- + Not reliant on *a priori* deciding the number of clusters
- Still relies on choices: Distance metric, linkage method, threshold





# Clustering horror story



UNIVERSITY  
OF OSLO

 Clustering horror story 

## Resting-state connectivity biomarkers define neurophysiological subtypes of depression

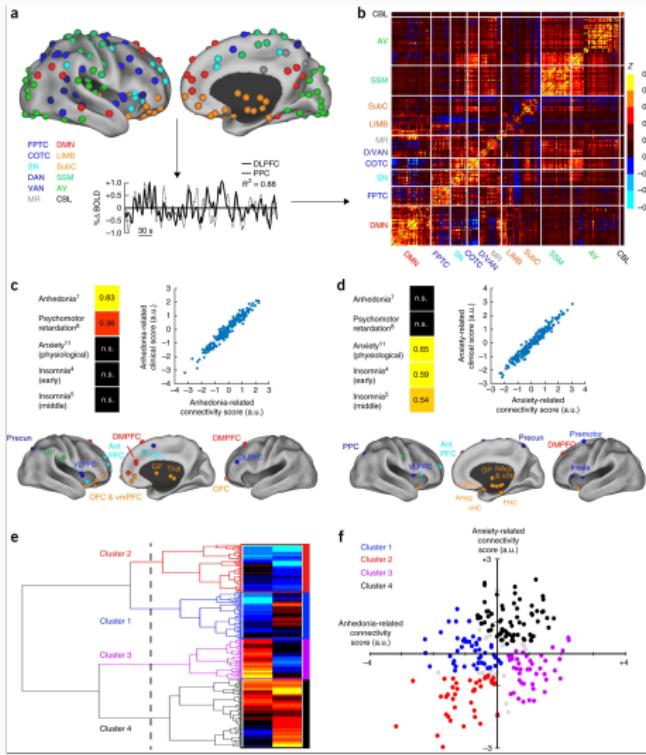
Andrew T Drysdale, Logan Grosenick, Jonathan Downar, Katharine Dunlop, Farrokh Mansouri, Yue Meng, Robert N Fetcho, Benjamin Zebley, Desmond J Oathes, Amit Etkin, Alan F Schatzberg, Keith Sudheimer, Jennifer Keller, Helen S Mayberg, Faith M Gunning, George S Alexopoulos, Michael D Fox, Alvaro Pascual-Leone, Henning U Voss, BJ Casey, Marc J Dubin & Conor Liston 

*Nature Medicine* 23, 28–38 (2017) | [Cite this article](#)

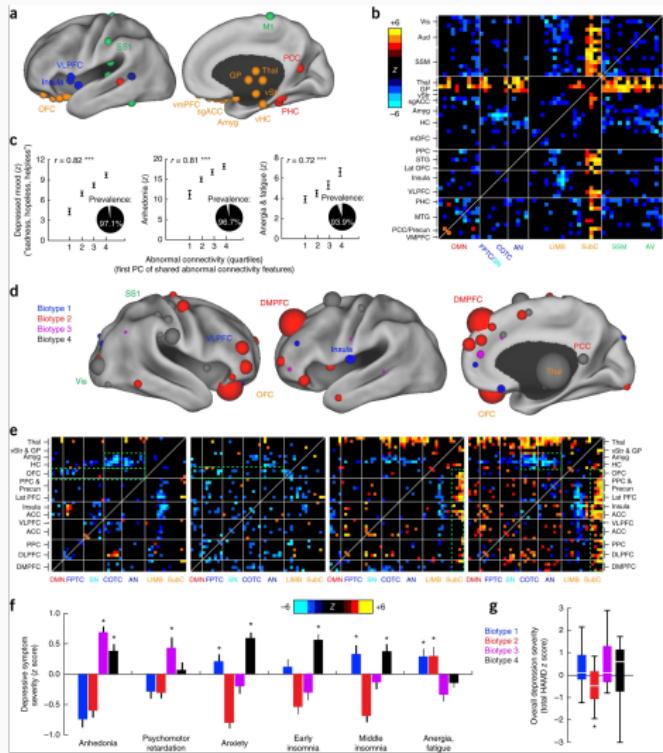
61k Accesses | 1348 Citations | 642 Altmetric | [Metrics](#)



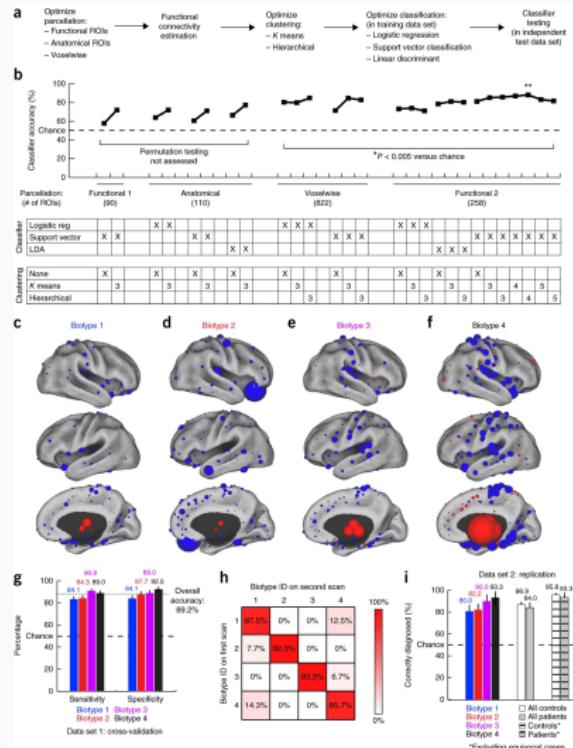
# Clustering horror story



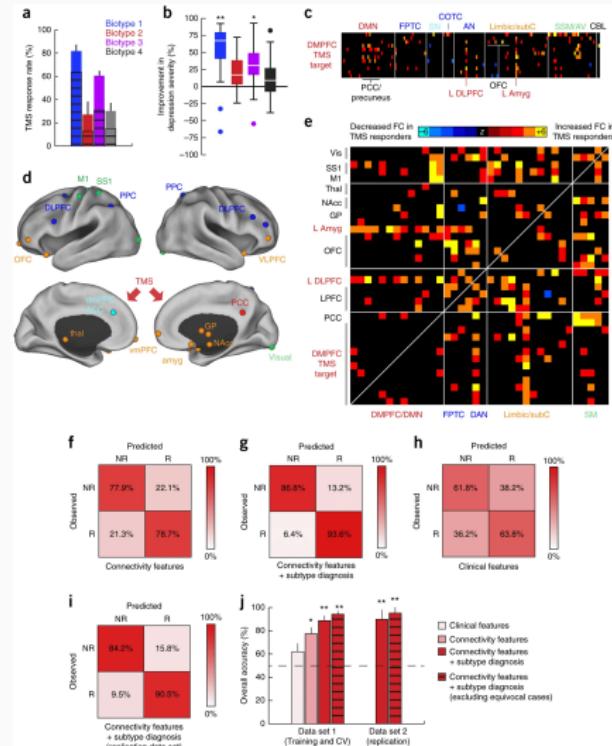
# Clustering horror story



# Clustering horror story



# Clustering horror story



# Clustering horror story



NeuroImage: Clinical  
Volume 22, 2019, 101796

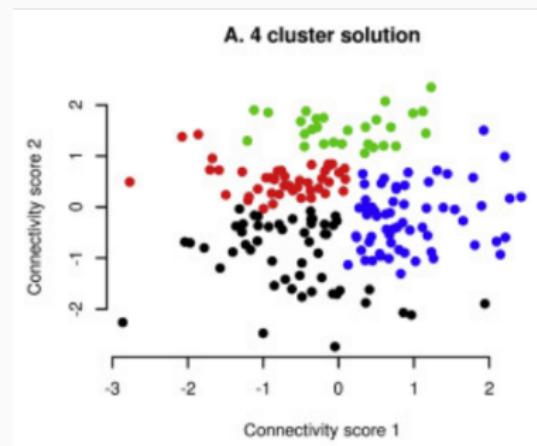


## Evaluating the evidence for biotypes of depression: Methodological replication and extension of Drysdale et al. (2017)

Richard Dinga <sup>a</sup>  , Lianne Schmaal <sup>b c</sup>, Brenda W.J.H. Penninx <sup>a</sup>, Marie Jose van Tol <sup>d</sup>,  
Dick J. Veltman <sup>a</sup>, Laura van Velzen <sup>a</sup>, Maarten Mennes <sup>e</sup>, Nic J.A. van der Wee <sup>f</sup>,  
Andre F. Marquand <sup>e</sup>

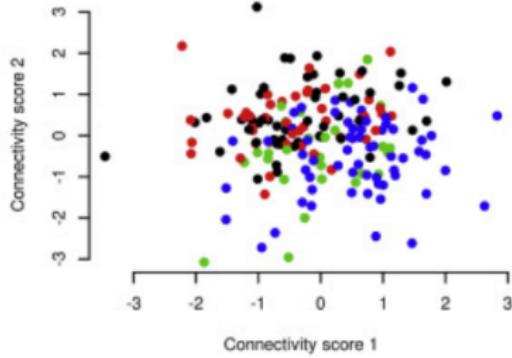


# Clustering horror story



# Clustering horror story

B. Cluster stability



# Clustering horror story

How can we avoid ending up with a clustering nightmare? 

- Quantitative evaluation of our clusters (see for instance Elements of Statistical Learning)
- Test cluster stability via cross-validation, half-split tests, bootstrap etc.
- Be cautious in interpretations



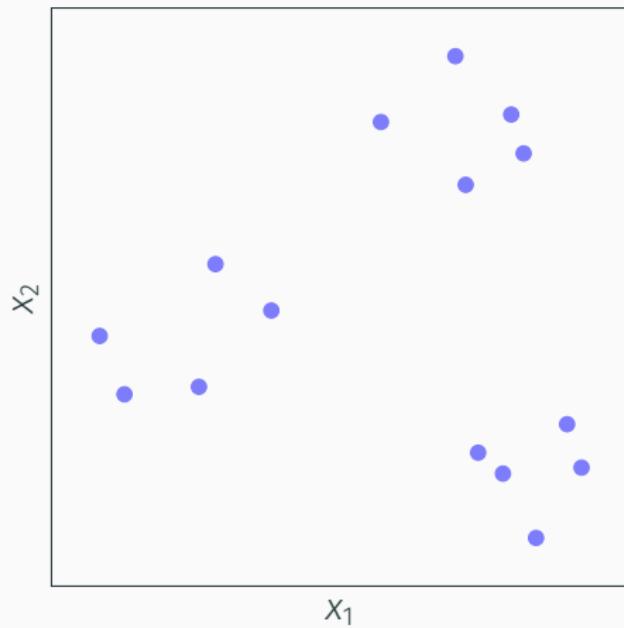
# Dimensionality reduction

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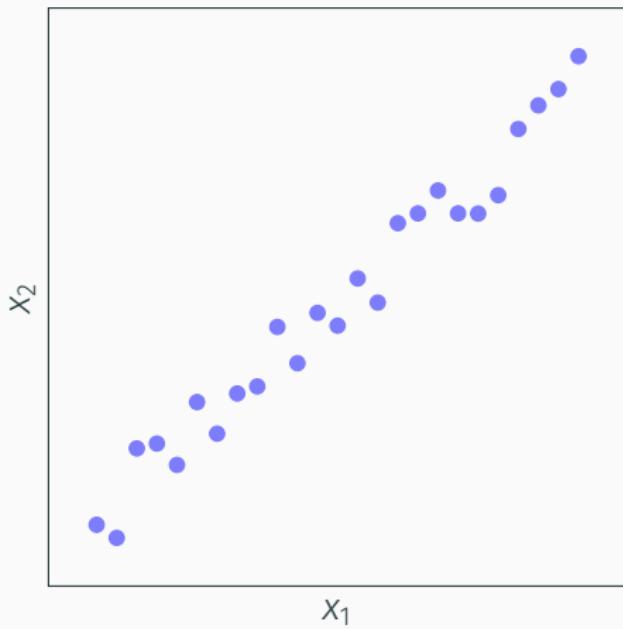


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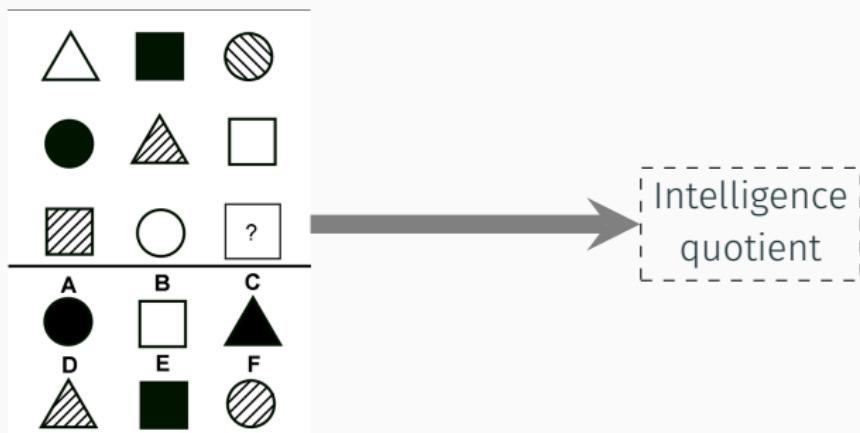
# Dimensionality reduction: Motivation



## Dimensionality reduction: Motivation



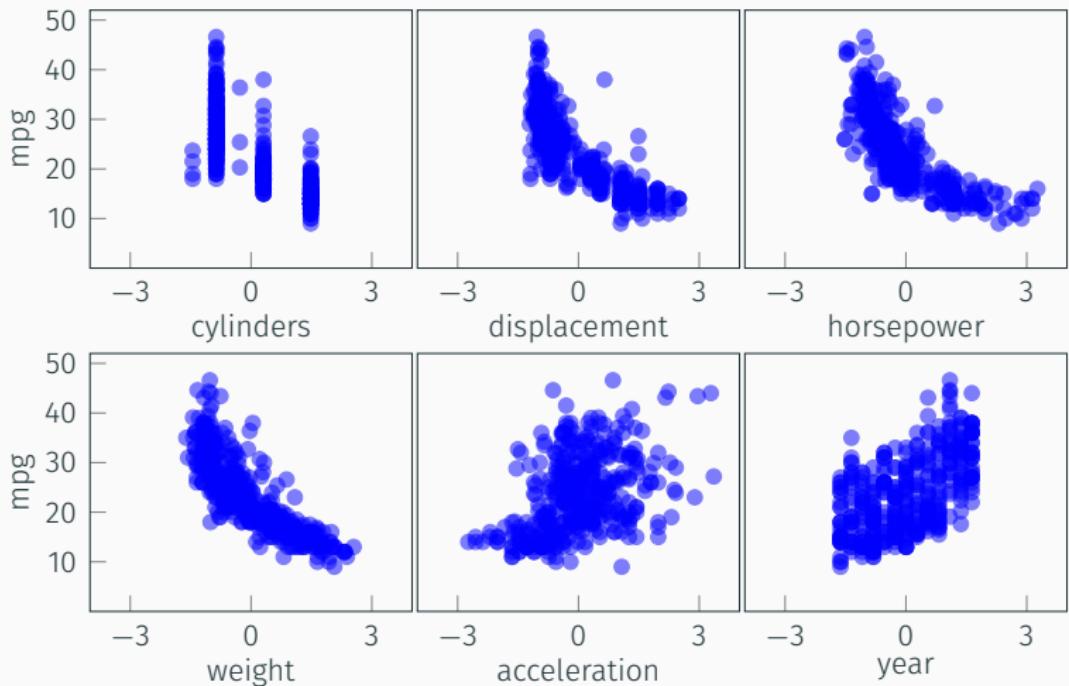
# Dimensionality reduction: Motivation



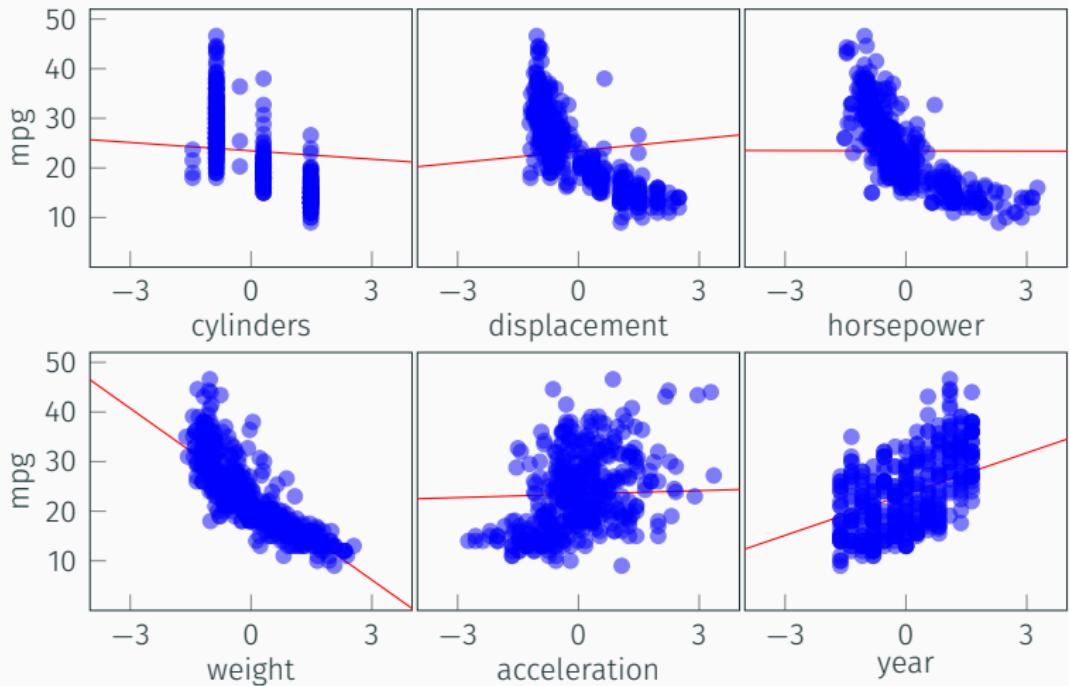
# Dimensionality reduction: Motivation



# Dimensionality reduction: Motivation



# Dimensionality reduction: Motivation



# Dimensionality reduction: Motivation

1	0.30	0.86	0.89	0.41	0.93
0.30	1	0.41	0.34	0.29	0.36
0.86	0.41	1	0.84	0.68	0.89
0.89	0.34	0.84	1	0.50	0.95
0.41	0.29	0.68	0.50	1	0.54
0.93	0.36	0.89	0.95	0.54	1

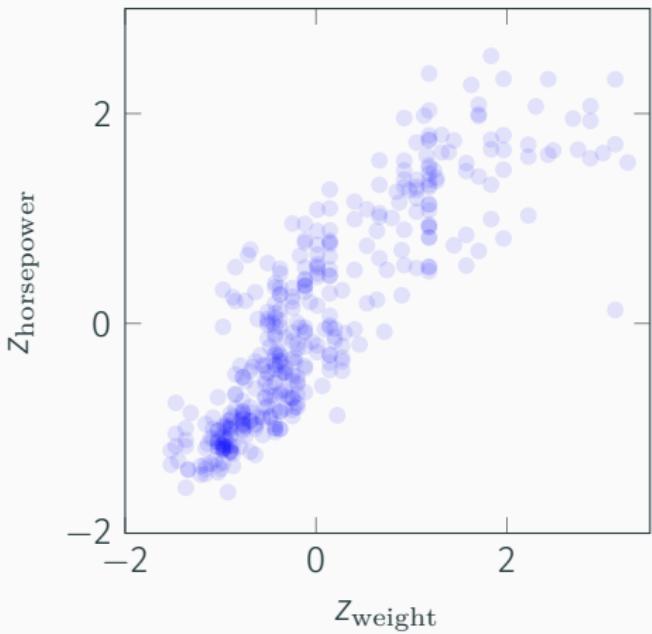


# Dimensionality reduction: Motivation

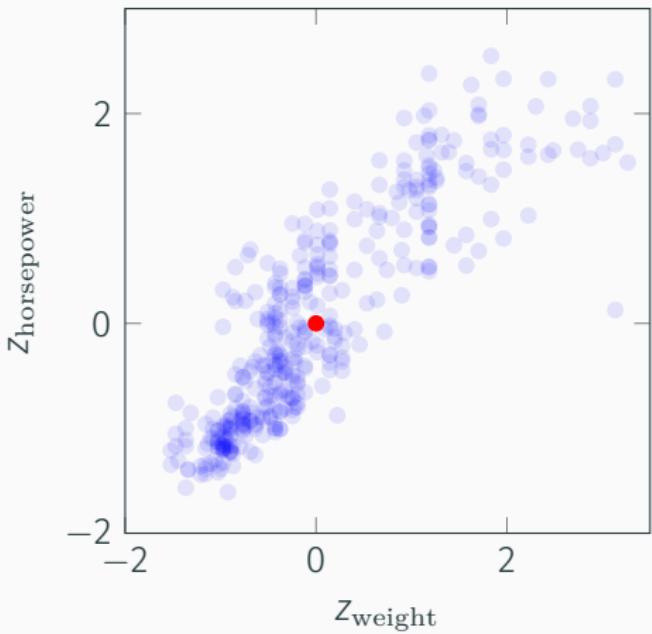
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0.30	1	0.41	0.34	0.29	0.36
0.86	0.41	1	0.84	0.68	0.89
0.89	0.34	0.84	1	0.50	0.95
0.41	0.29	0.68	0.50	1	0.54
0.93	0.36	0.89	0.95	0.54	1



# Dimensionality reduction: Principal component analysis



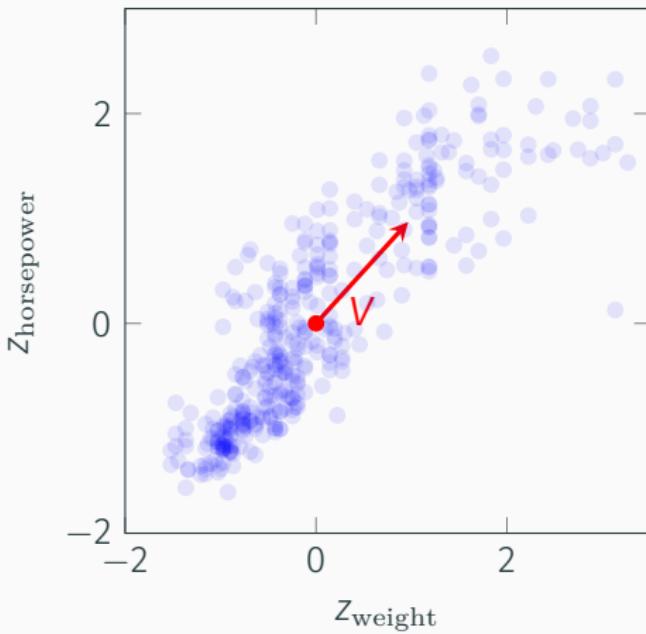
# Dimensionality reduction: Principal component analysis



$c \rightarrow$  center of the data



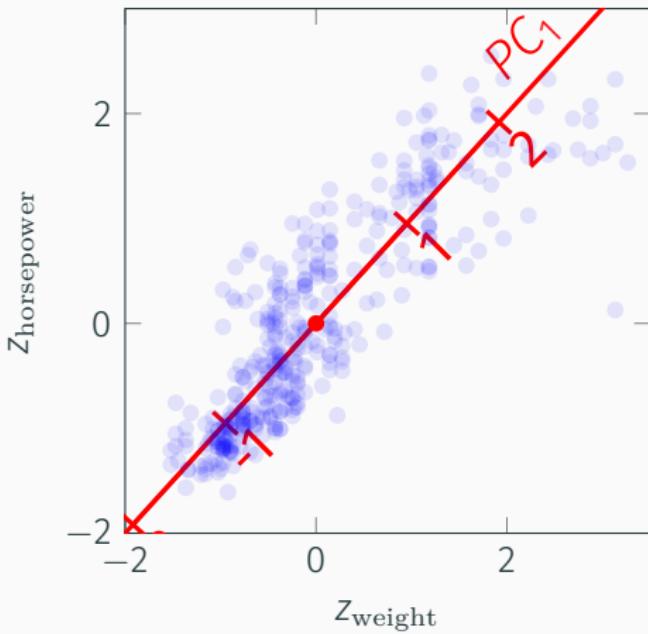
# Dimensionality reduction: Principal component analysis



$v \rightarrow$  direction of maximum variance



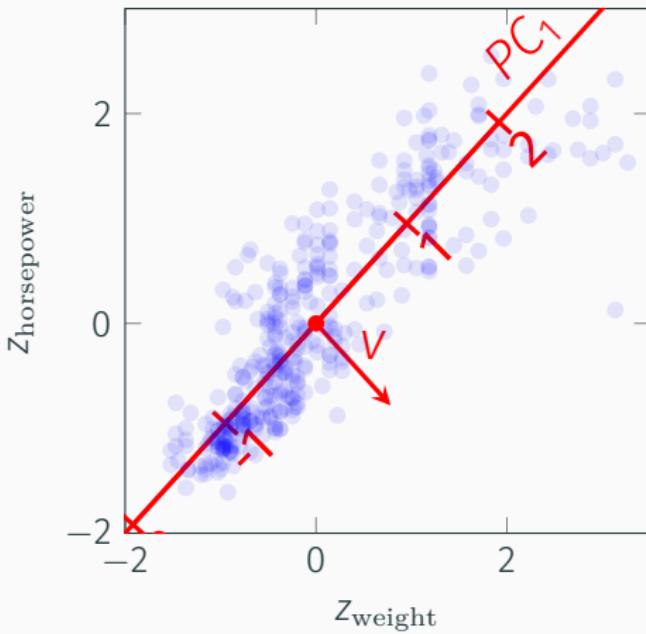
# Dimensionality reduction: Principal component analysis



$$PC_1 \rightarrow 0.69 * Z_{\text{horsepower}} + 0.71 * Z_{\text{weight}}$$



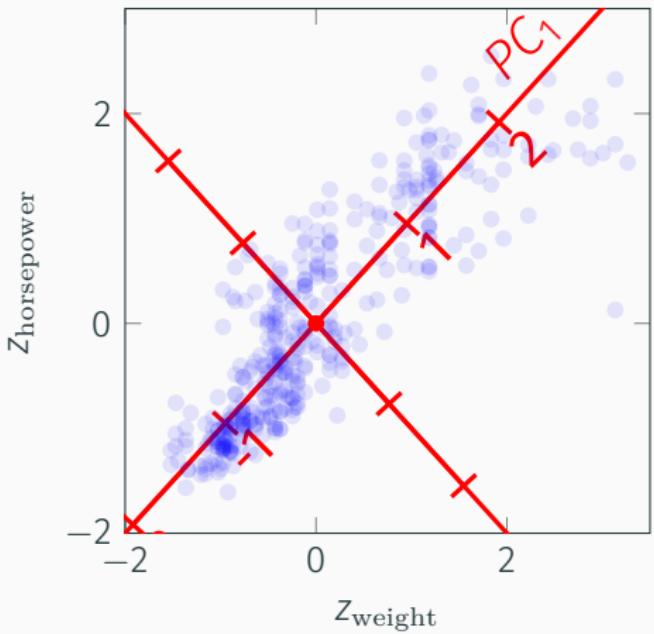
# Dimensionality reduction: Principal component analysis



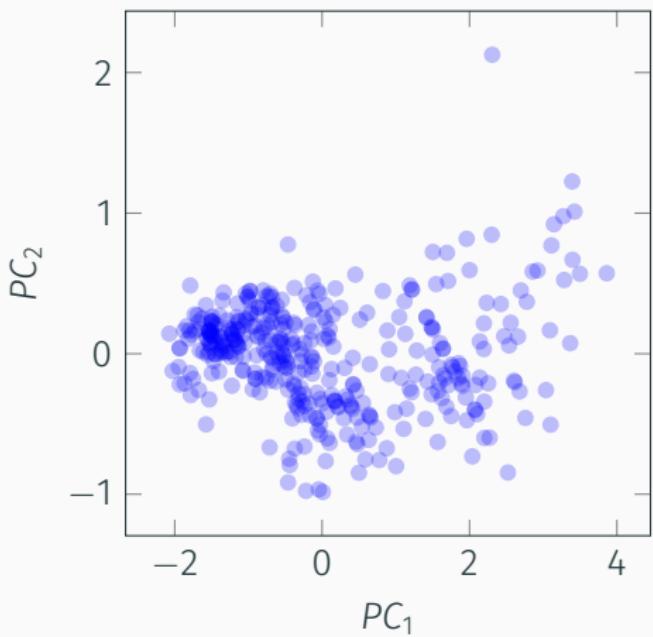
$v \rightarrow$  direction of maximum variance **orthogonal** to  $PC_1$



# Dimensionality reduction: Principal component analysis



# Dimensionality reduction: Principal component analysis



# Dimensionality reduction: Principal component analysis

mpg	horsepower	weight	PC1	PC2
18	130	3504	0.908	0.303
15	165	3693	1.709	0.517
18	150	3436	1.219	0.455
16	150	3433	1.217	0.457
17	140	3449	1.046	0.260
15	198	4341	2.856	0.583
14	220	4354	3.272	0.977



# Dimensionality reduction: Principal component analysis

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$$PC_1 = 0.69 * z_{\text{horsepower}} + 0.71 * z_{\text{weight}}$$

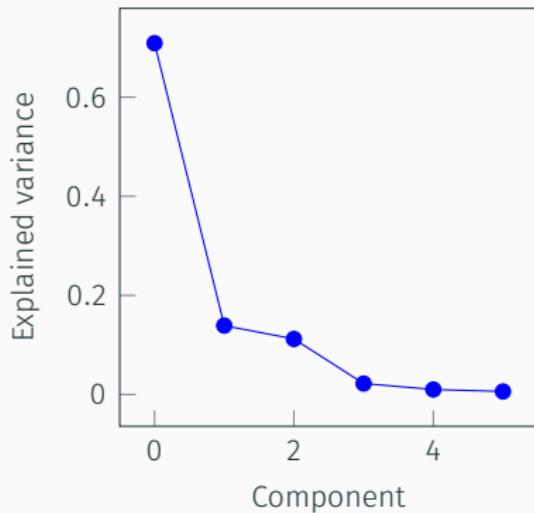


# Dimensionality reduction: Principal component analysis

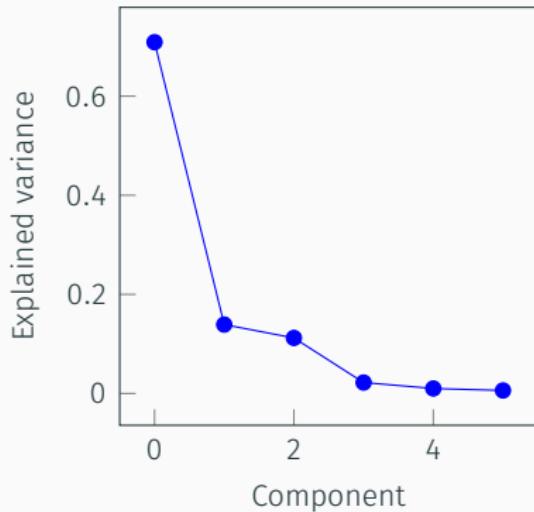
<http://localhost:8888/notebooks/notebooks%2FPCA.ipynb>



# Dimensionality reduction: Principal component analysis



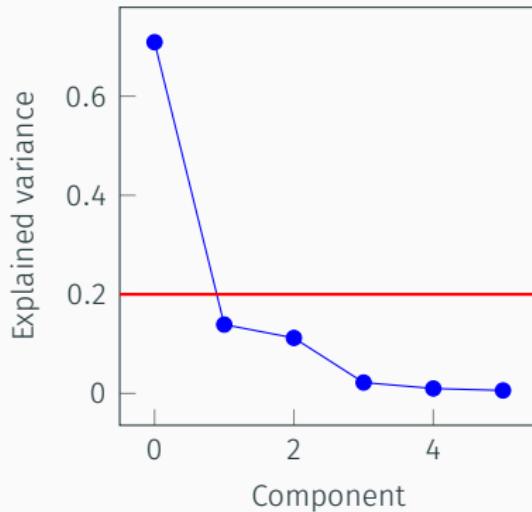
# Dimensionality reduction: Principal component analysis



$$\hat{y} = \beta_0 + \sum_{i=0}^n \beta_i PC_i$$



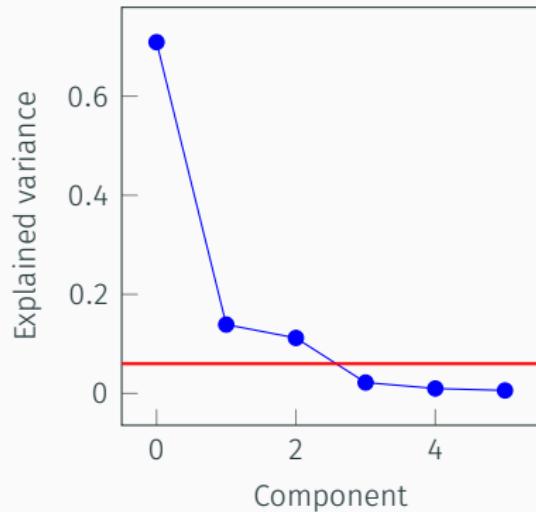
# Dimensionality reduction: Principal component analysis



$$\hat{y} = \beta_0 + \sum_{i=1}^1 \beta_i PC_i$$



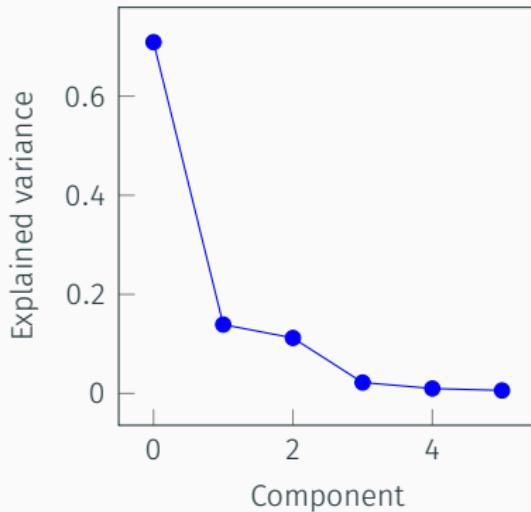
# Dimensionality reduction: Principal component analysis



$$\hat{y} = \beta_0 + \sum_{i=1}^3 \beta_i PC_i$$



# Dimensionality reduction: Principal component analysis



$$\hat{y} = \beta_0 + \sum_{i=0}^n \beta_i PC_i$$

$n$  decided via a validation set, tested in a **held-out test set**



# Dimensionality reduction: Principal component analysis

Principal component analysis: Transforms our dataset by computing *principal components* to replace our original variables.

- Principal components are:
  - Linear combinations of the original variables
  - Orthogonal to each other, meaning that they capture different signals in our data (linearly uncorrelated)
- They can be useful for:
  - (Qualitatively) understanding the signal in our data
  - Reducing the number of predictors for modelling



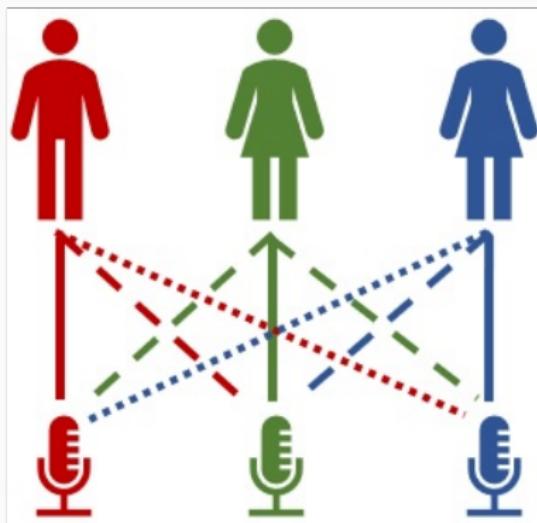
# Dimensionality reduction: Principal component analysis

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  - Reducing the number of predictors for modelling



# Dimensionality reduction: Independent component analysis



# Dimensionality reduction: Independent component analysis

Principal component analysis: Create orthogonal components that are linear combinations of our variables:

$$PC_0 = \beta_0x_0 + \beta_1x_1 + \dots + \beta_nx_n,$$

$$PC_1 = \gamma_0x_0 + \gamma_1x_1 + \dots + \gamma_nx_n,$$

$PC_0 \perp PC_1$  (e.g. no linear correlation)

Independent component analysis: Represent each of our original variables as a linear combination of underlying sources:

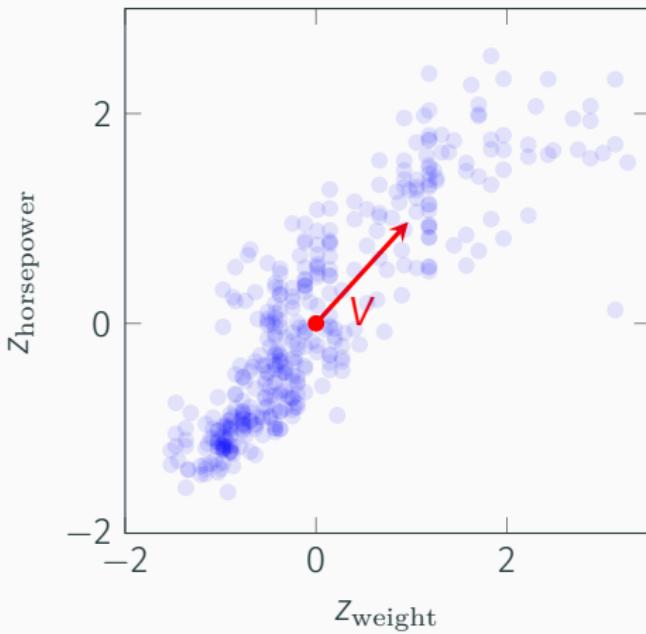
$$x_0 = \alpha_{00}s_0 + \alpha_{01}s_1 + \dots + \alpha_{0n}s_n,$$

$$x_1 = \alpha_{10}s_0 + \alpha_{11}s_1 + \dots + \alpha_{1n}s_n,$$

$s_0 \perp\!\!\!\perp s_1$  (e.g. statistically independent)



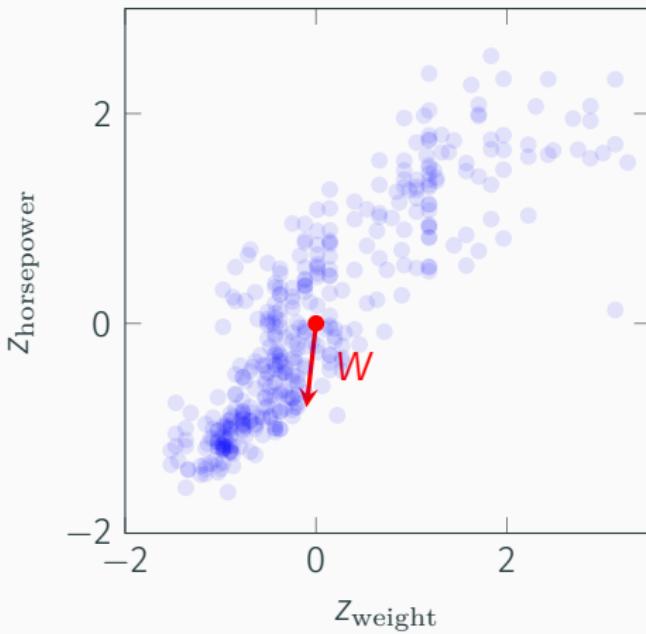
# Dimensionality reduction: Independent component analysis



$v \rightarrow$  direction of greatest variance



# Dimensionality reduction: Independent component analysis



$w \rightarrow$  direction that maximizes the non-Gaussianity of  $w^T X$



# Dimensionality reduction: Independent component analysis

[https://scikit-learn.org/dev/auto\\_examples/decomposition/plot\\_ica\\_blind\\_source\\_separation.html#sphx-glr-auto-examples-decomposition-plot-ica-blind-source-separation-py](https://scikit-learn.org/dev/auto_examples/decomposition/plot_ica_blind_source_separation.html#sphx-glr-auto-examples-decomposition-plot-ica-blind-source-separation-py)

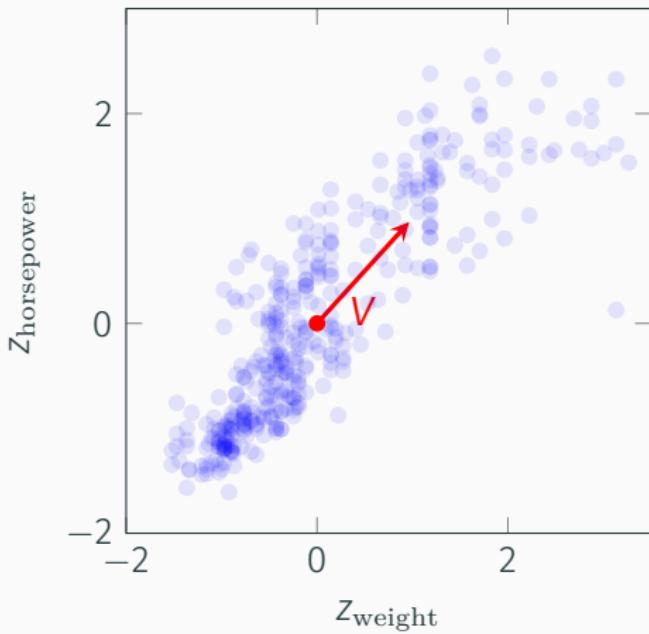


# Dimensionality reduction: Independent component analysis

<https://PMC7162660>



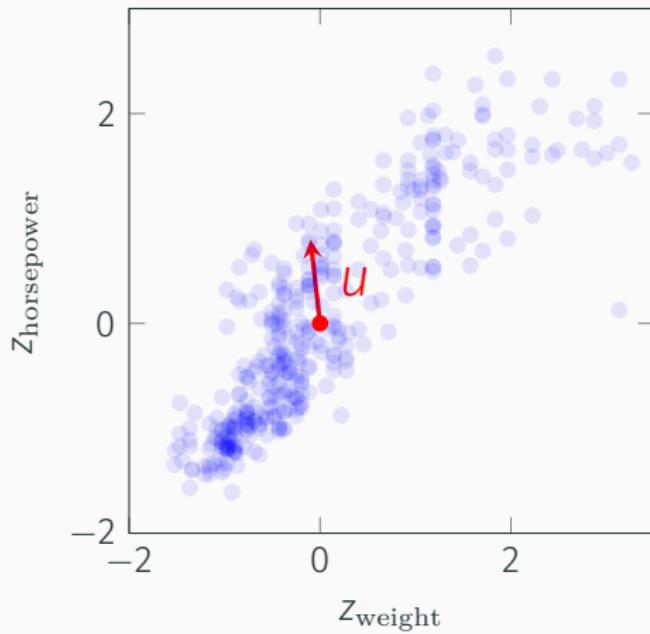
# Dimensionality reduction: Partial least squares



$v \rightarrow$  direction of maximum variance



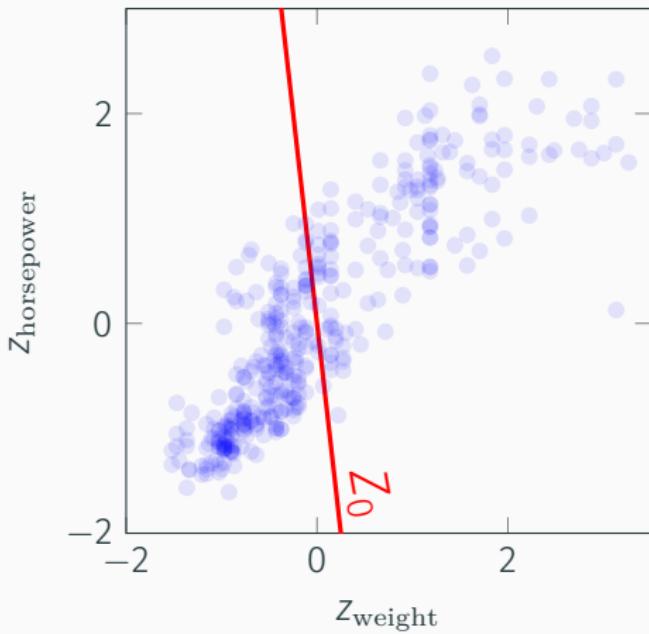
# Dimensionality reduction: Partial least squares



$u \rightarrow$  direction of greatest covariance between  $X$  and  $y$



# Dimensionality reduction: Partial least squares



# Dimensionality reduction: Partial least squares

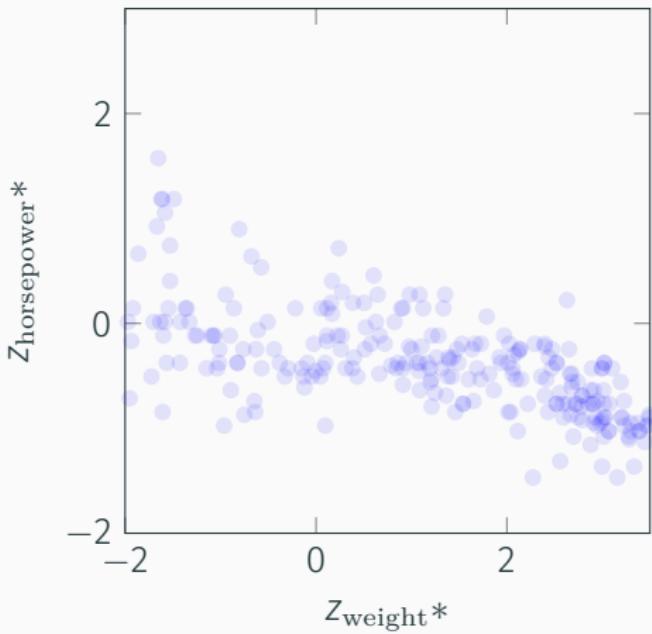
$$z_{\text{horsepower}*} = z_{\text{horsepower}} - Z_0$$

$$z_{\text{weight}*} = z_{\text{weight}} - Z_0$$

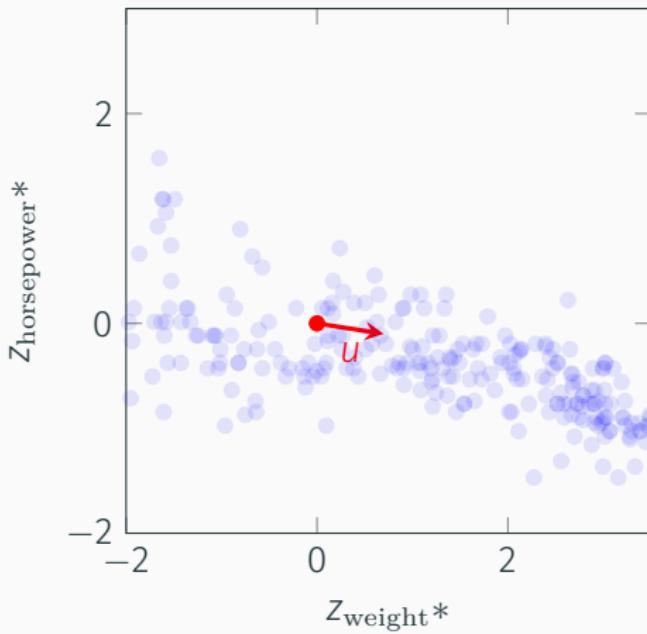
$$y* = y - Z_0$$



# Dimensionality reduction: Partial least squares



# Dimensionality reduction: Partial least squares



$u \rightarrow$  direction of greatest covariance between  $X^*$  and  $y^*$



# Dimensionality reduction: Partial least squares

Principal component analysis: Create orthogonal components that are linear combinations of our variables:

$$PC_0 = \beta_0x_0 + \beta_1x_1 + \dots + \beta_nx_n,$$

$$PC_1 = \gamma_0x_0 + \gamma_1x_1 + \dots + \gamma_nx_n,$$

$PC_0 \perp PC_1$  (e.g. no linear correlation),

that maximize the variance of  $X$

Partial least squares: Create orthogonal components that are linear combinations of our variables:

$$Z_0 = \beta_0x_0 + \beta_1x_1 + \dots + \beta_nx_n,$$

$$Z_1 = \gamma_0x_0 + \gamma_1x_1 + \dots + \gamma_nx_n,$$

$Z_0 \perp PC_1$  (e.g. no linear correlation),

that maximize the covariance between  $X$  and  $y$



# Dimensionality reduction: Summary

Dimensionality reduction techniques allow us to reduce the number of variables in our dataset to either aid interpretation, or improve our models through implicit regularization.

- Principal component analysis (PCA): Finds components that are orthogonal and maximize variance
- Independent component analysis (ICA): Finds components that are non-Gaussian and statistically independent
- Partial least squares (PLS): Finds components that are orthogonal and maximize covariance between  $X$  and  $y$

