

# Unsupervised Learning 2

# Representation Learning

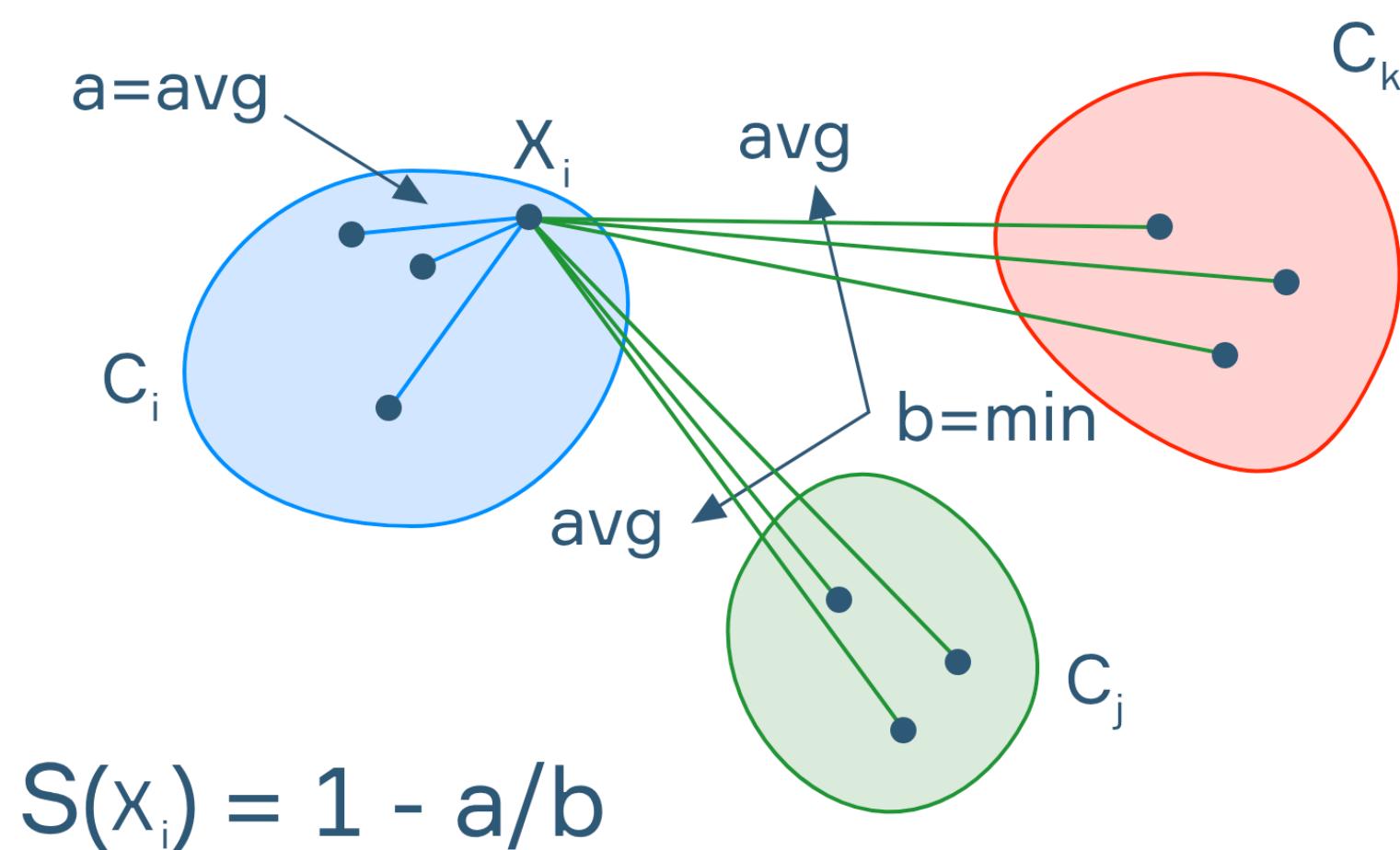
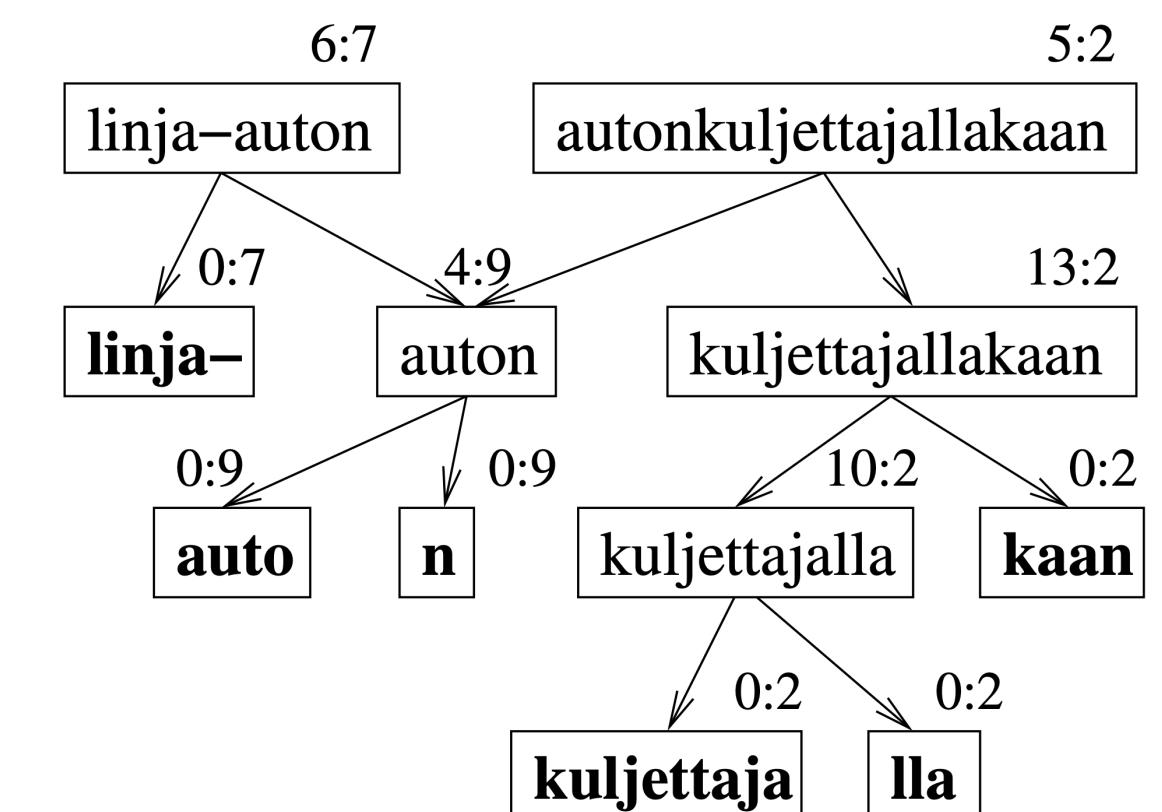
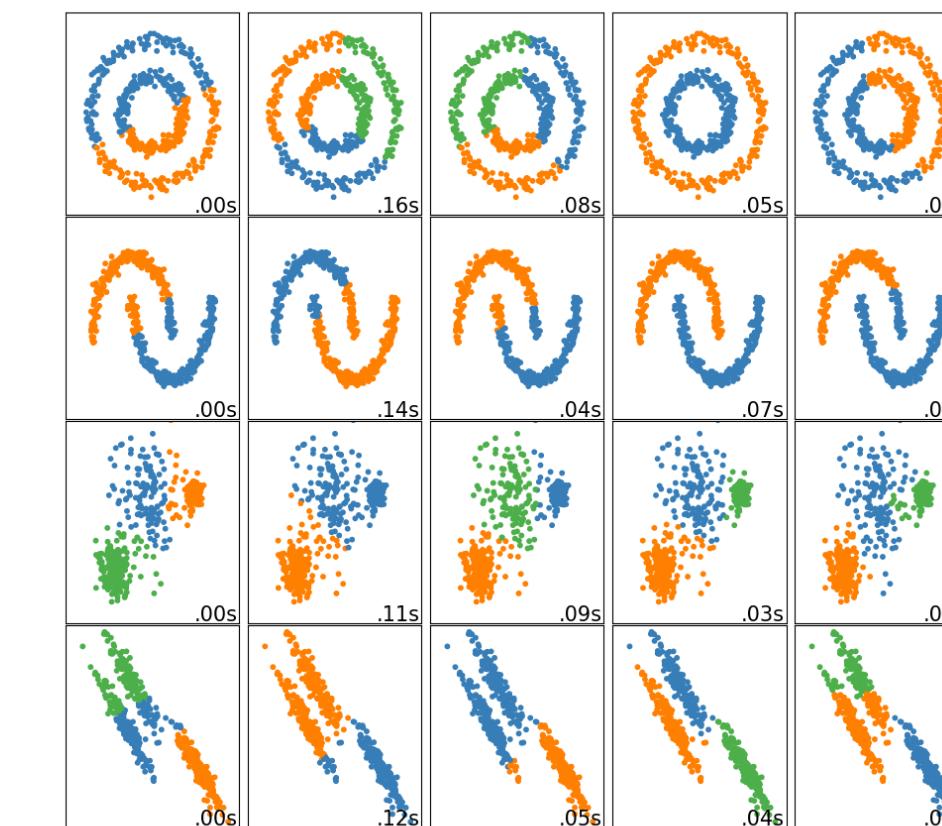
DSCC 251/451: Machine Learning with Limited Data

C.M. Downey

Spring 2026

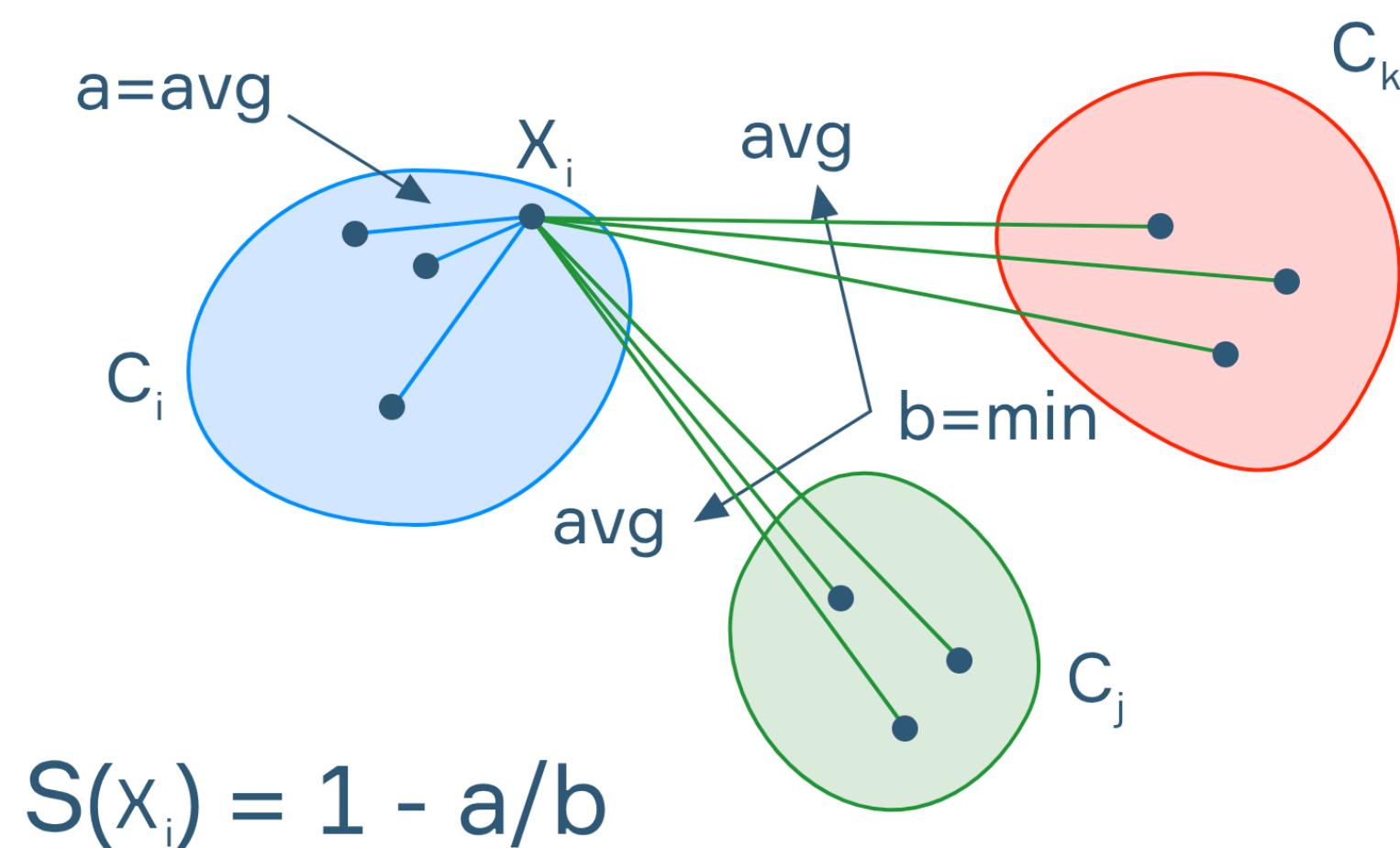
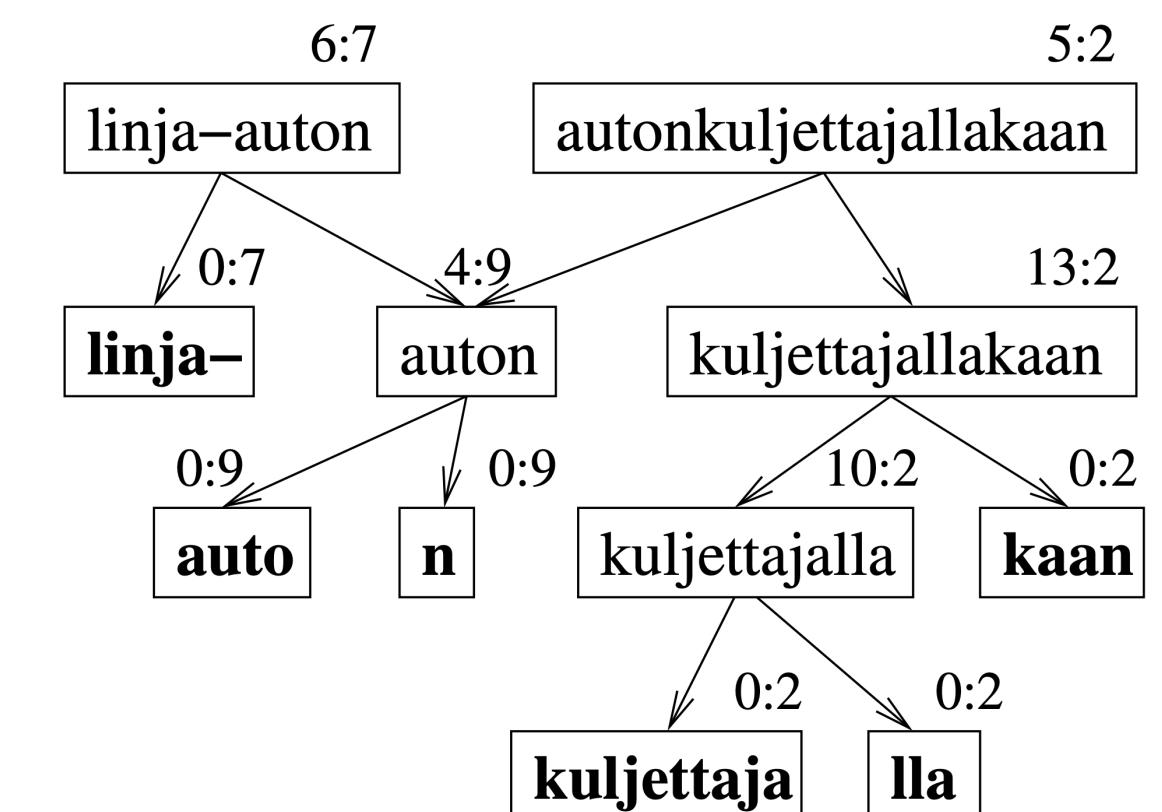
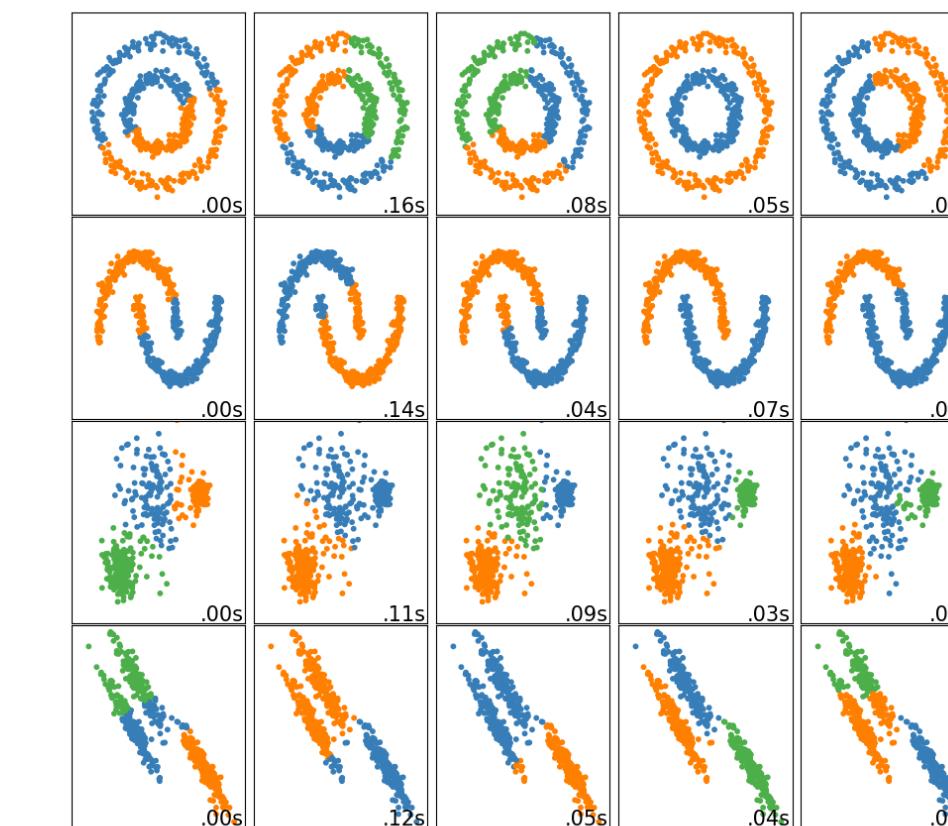
# Recap and Roadmap

# Recap



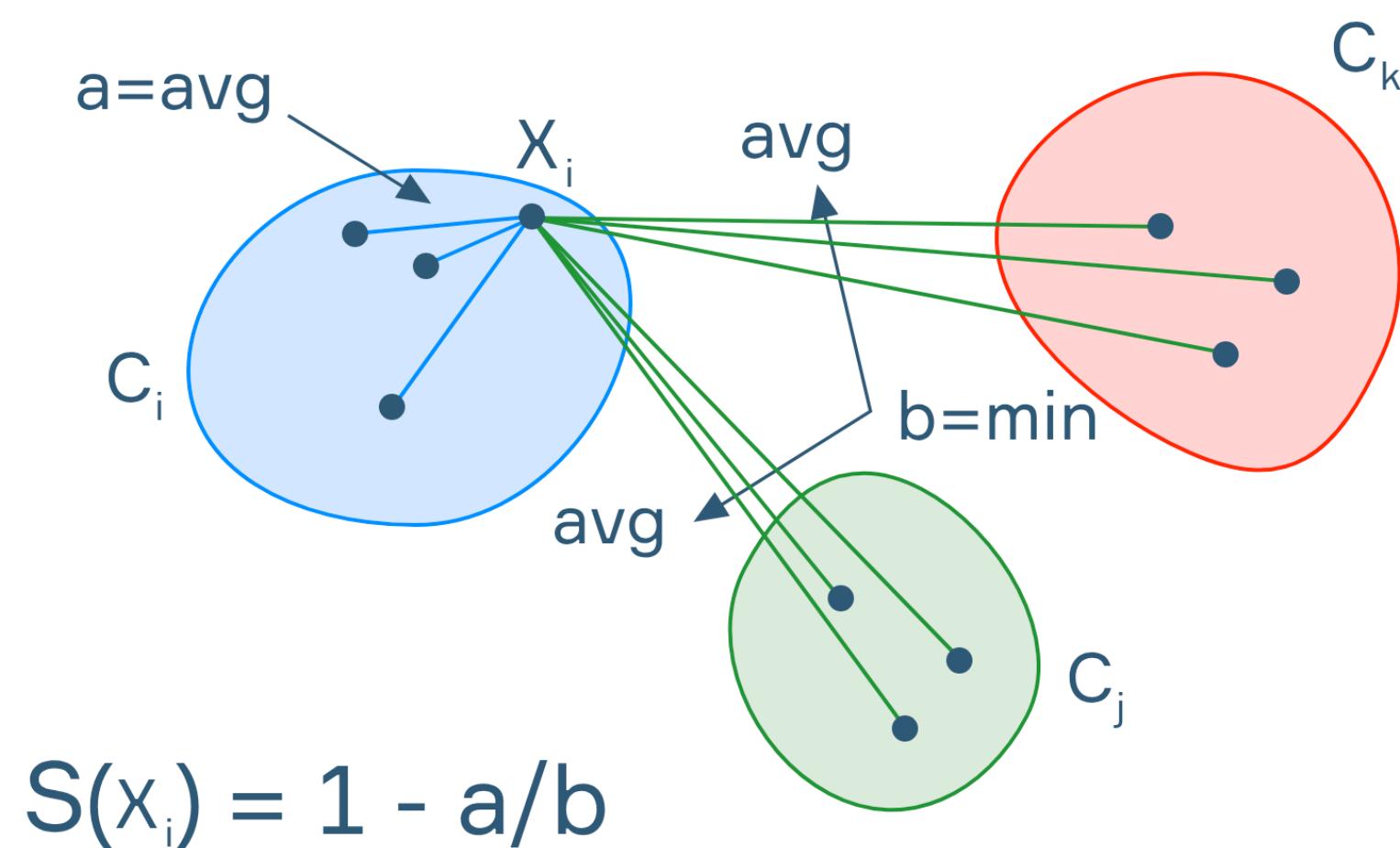
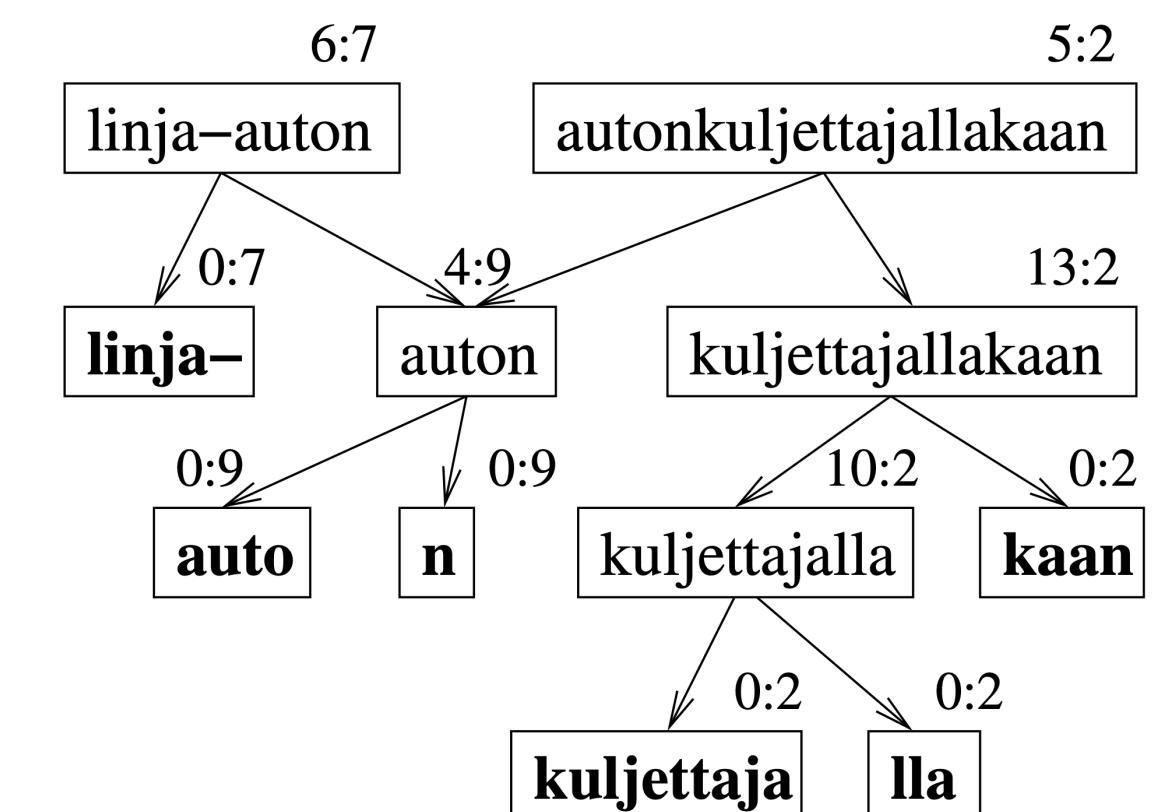
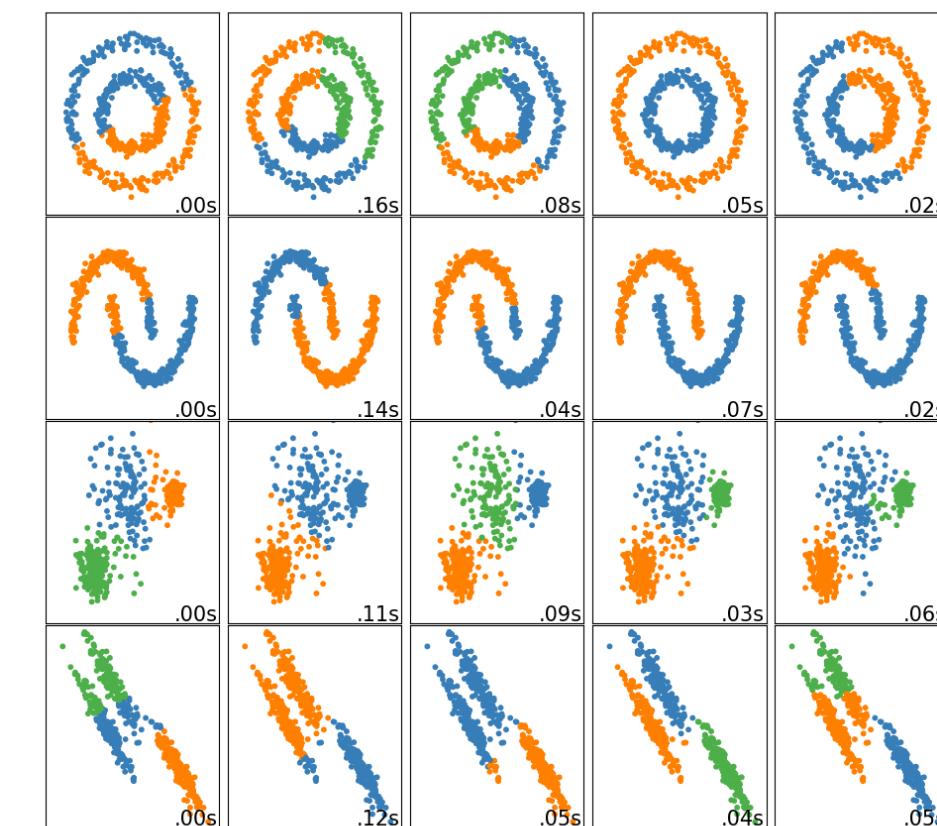
# Recap

- Unsupervised Learning = **discovering structure in (unlabeled) data**



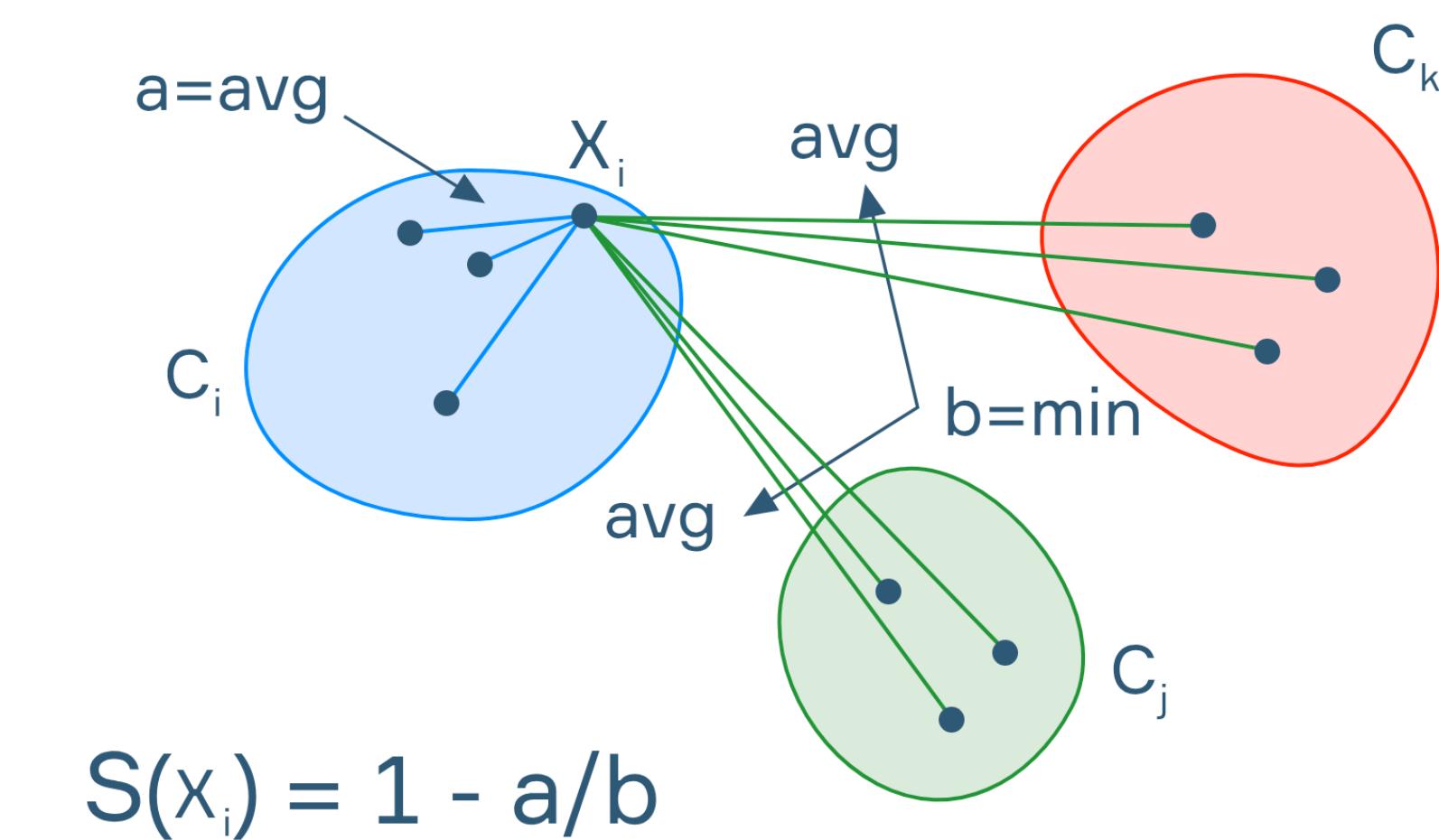
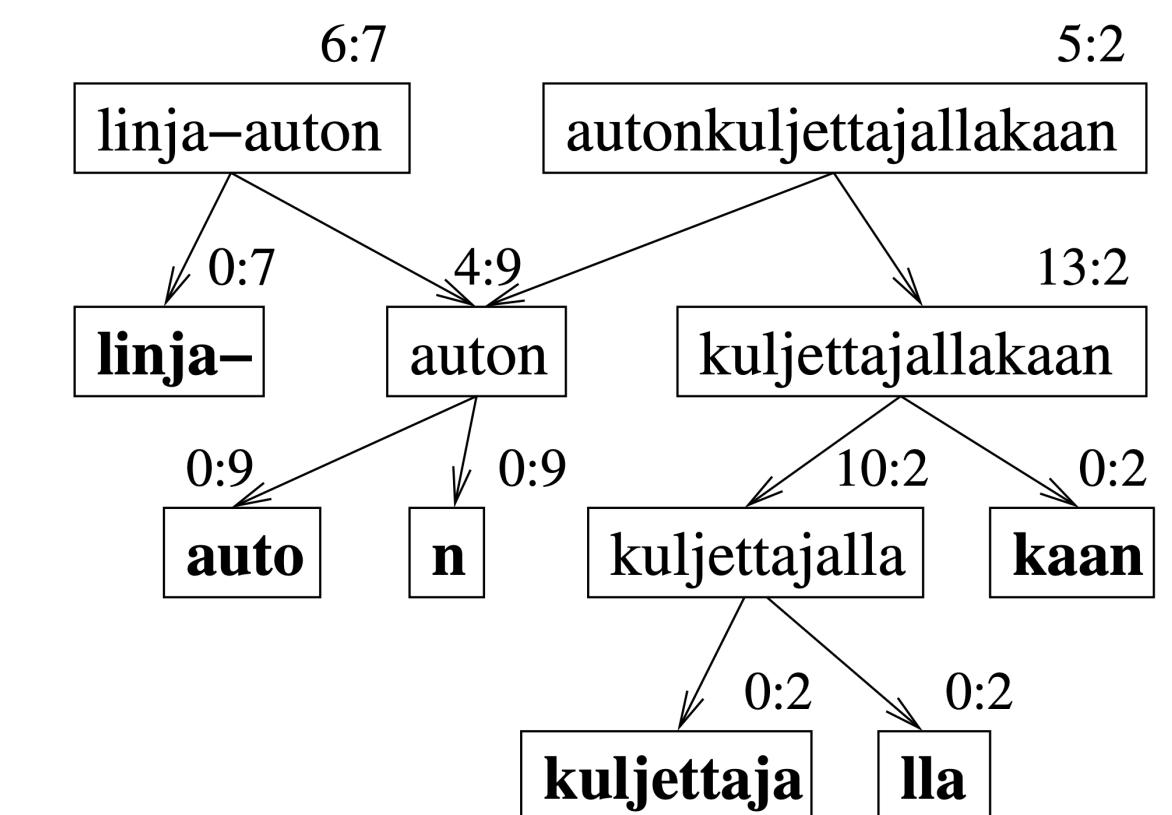
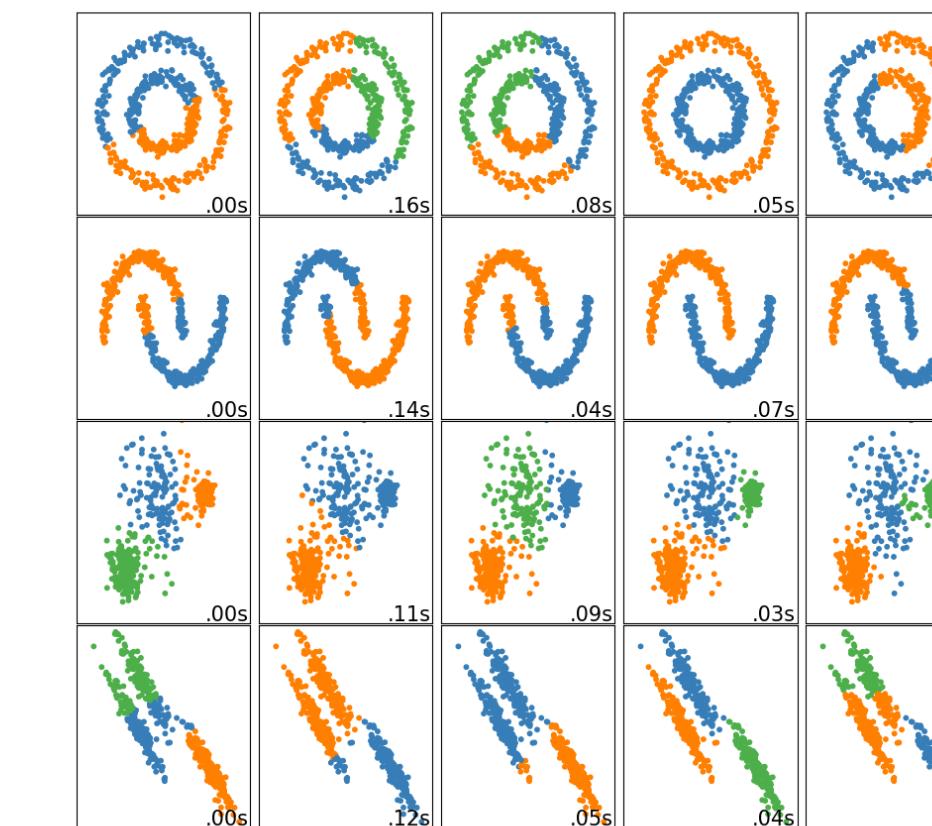
# Recap

- Unsupervised Learning = **discovering structure in (unlabeled) data**
- Usually have to optimize a **surrogate objective**, since you can't optimize a supervised one
  - Hope/hypothesis: this **correlates** with something you care about

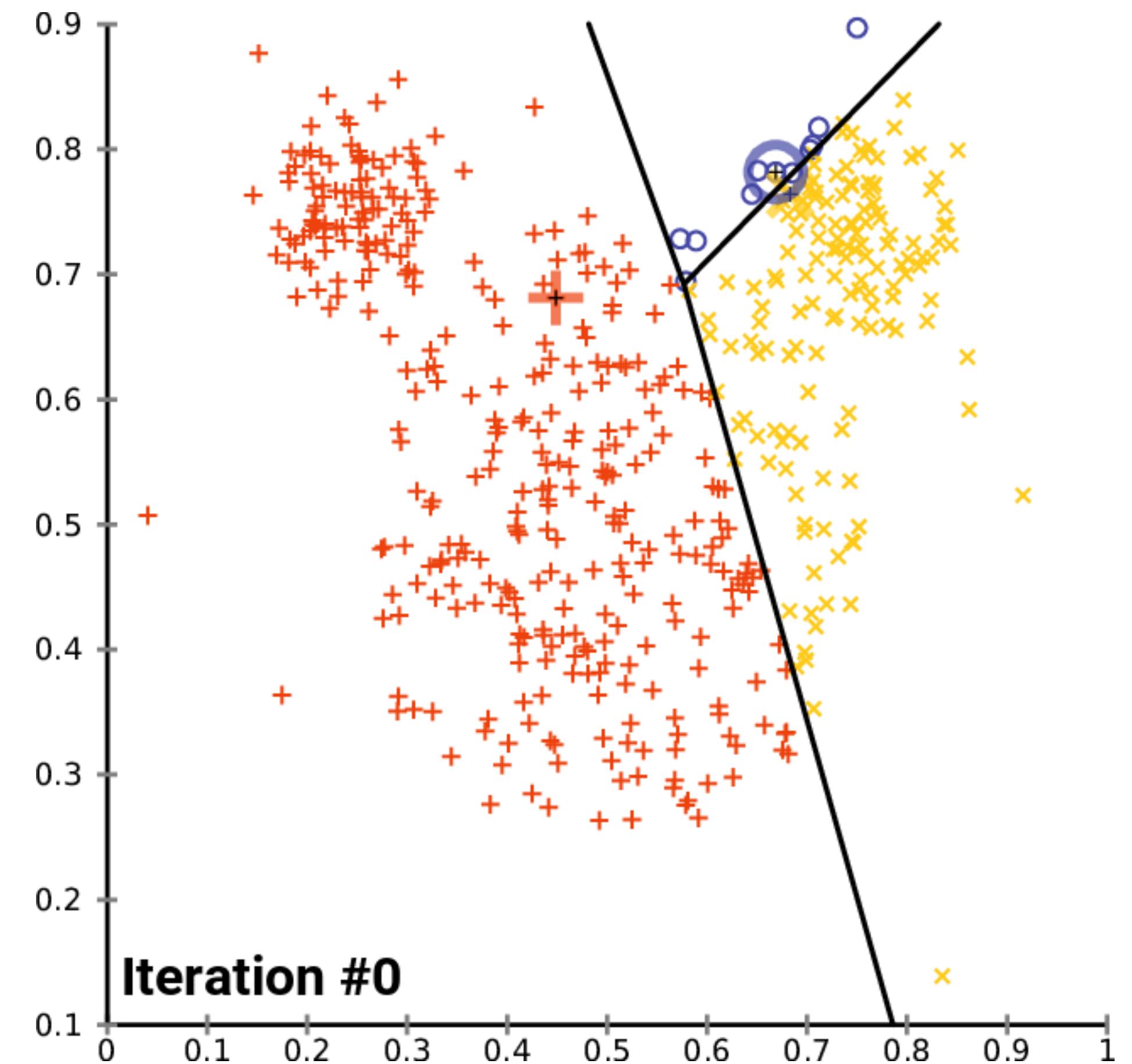


# Recap

- Unsupervised Learning = **discovering structure in (unlabeled) data**
- Usually have to optimize a **surrogate objective**, since you can't optimize a supervised one
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- Last time: discovering **discrete structure** (clusters, segmentation)
  - Hope these stand in for **supervised outputs/labels**

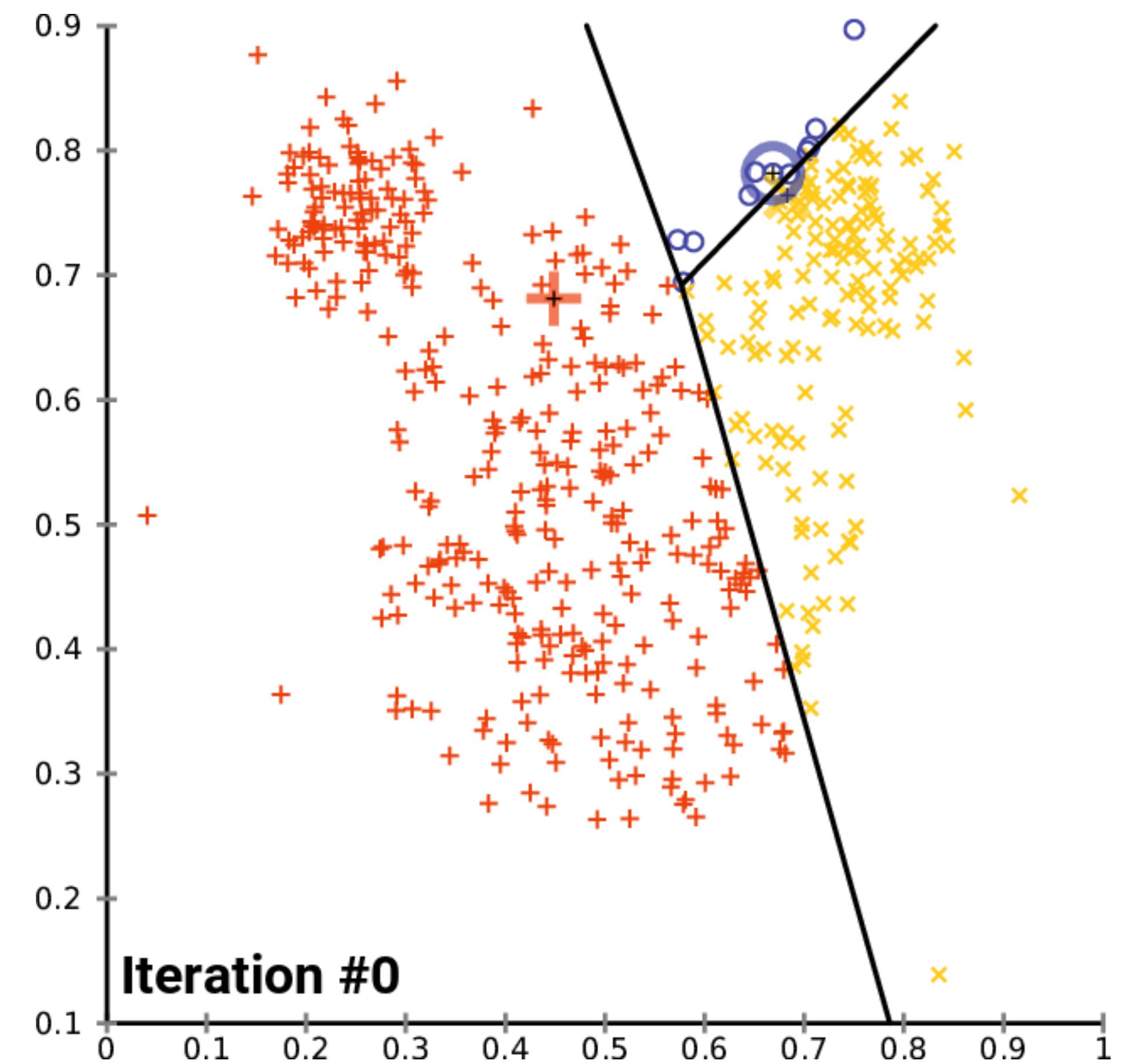


# Clustering



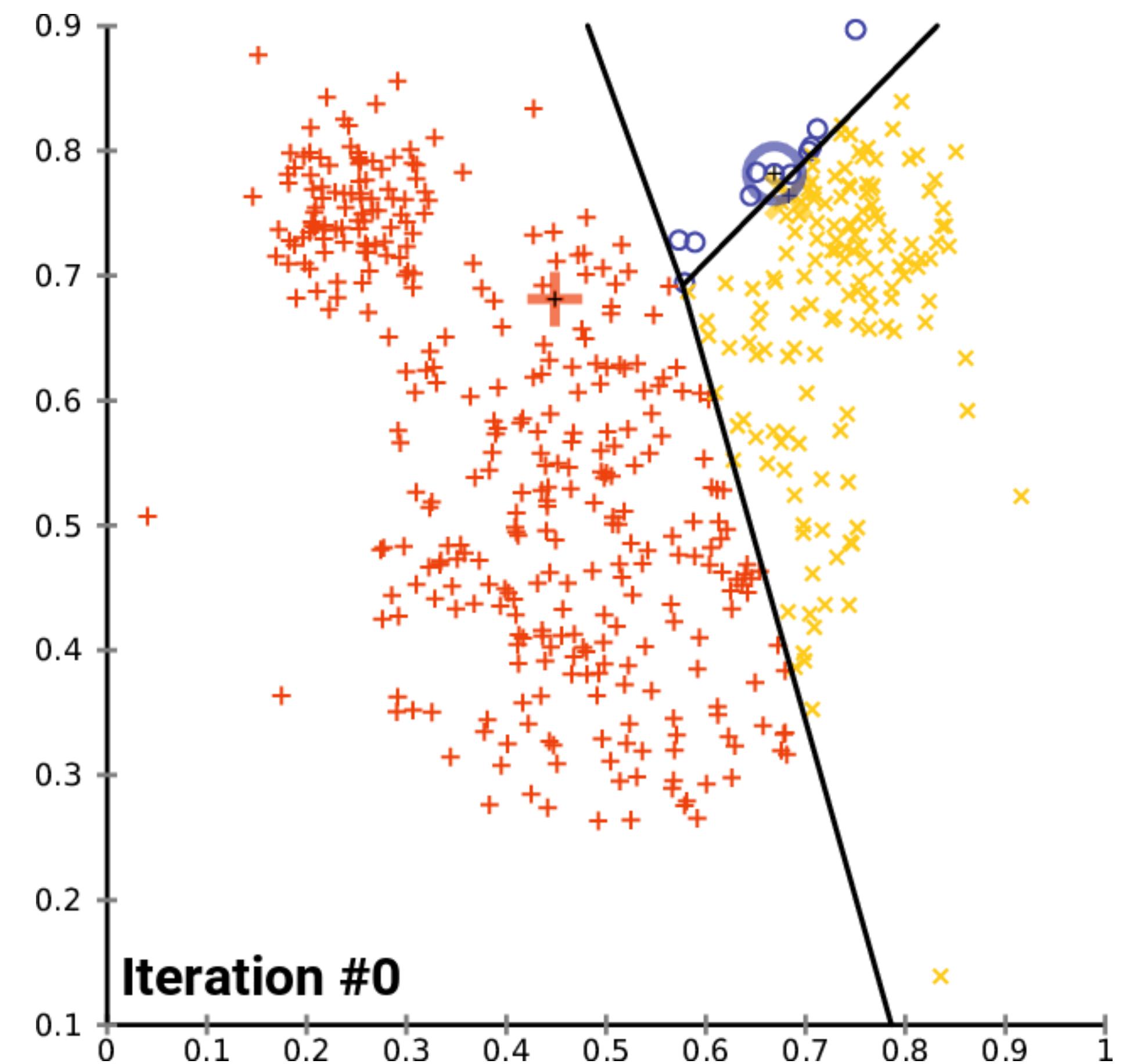
# Clustering

- Goal: find **groupings** in data



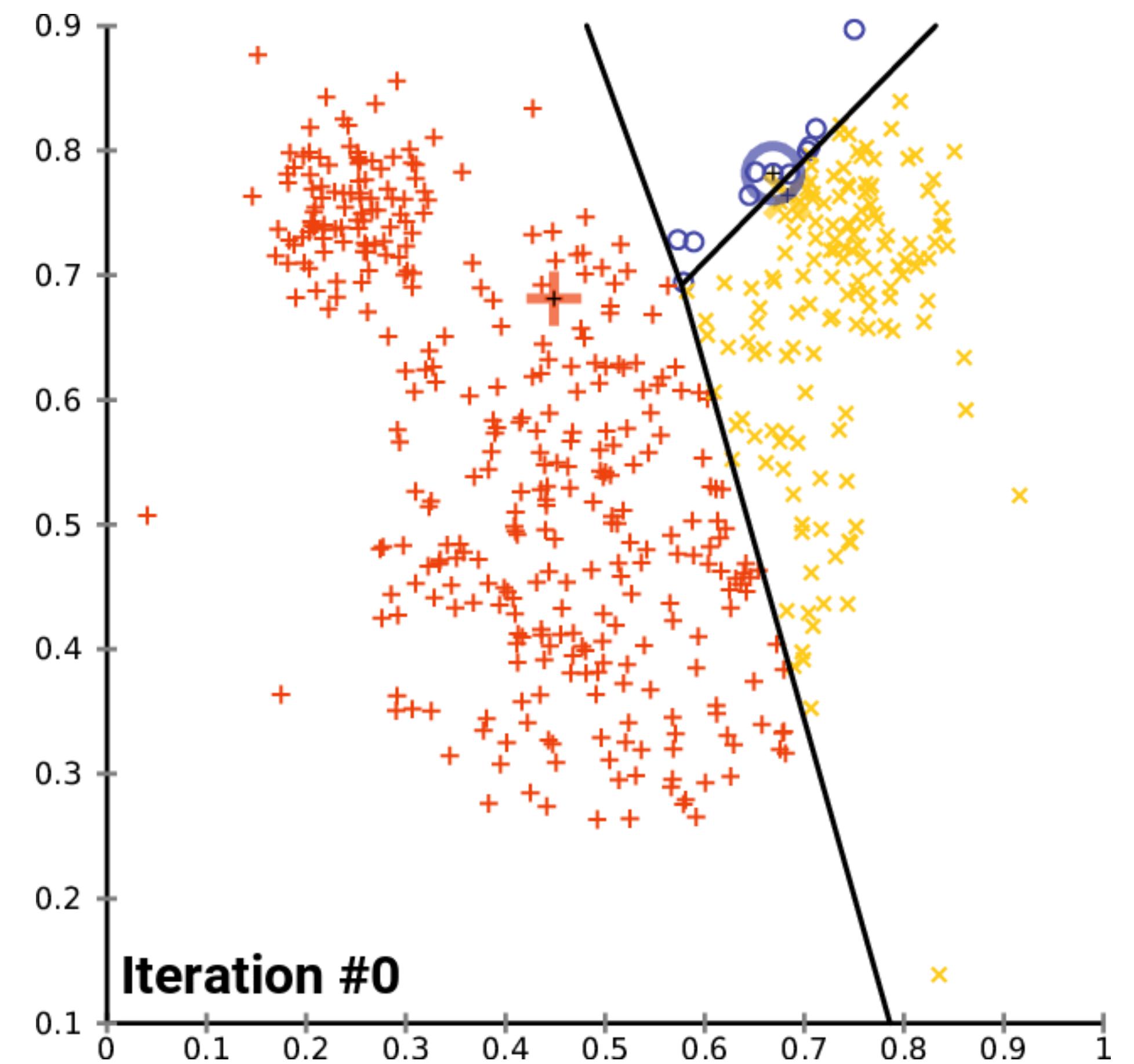
# Clustering

- Goal: find **groupings** in data
- Surrogate metric: **cluster quality**
  - K-means: minimize **Within-Cluster Sum of Squares** (WCSS, "compactness")
  - Silhouette Score: measures **coherence** and **distinctness**



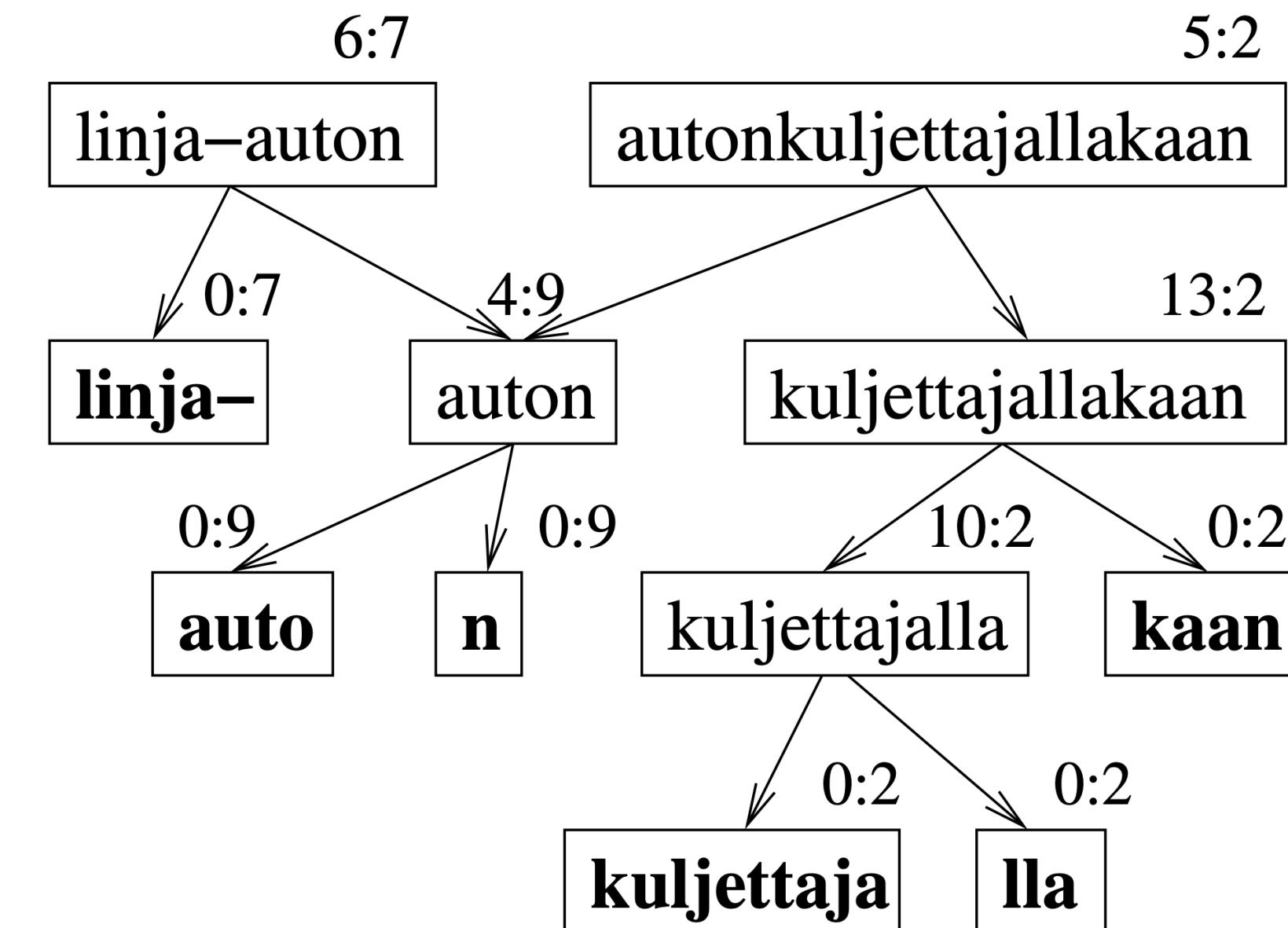
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  - Silhouette Score: measures **coherence** and **distinctness**
- Inductive biases: all kinds!
  - K-means biased towards **spherical clusters**
  - Other algorithms have different biases



# Sequence Segmentation

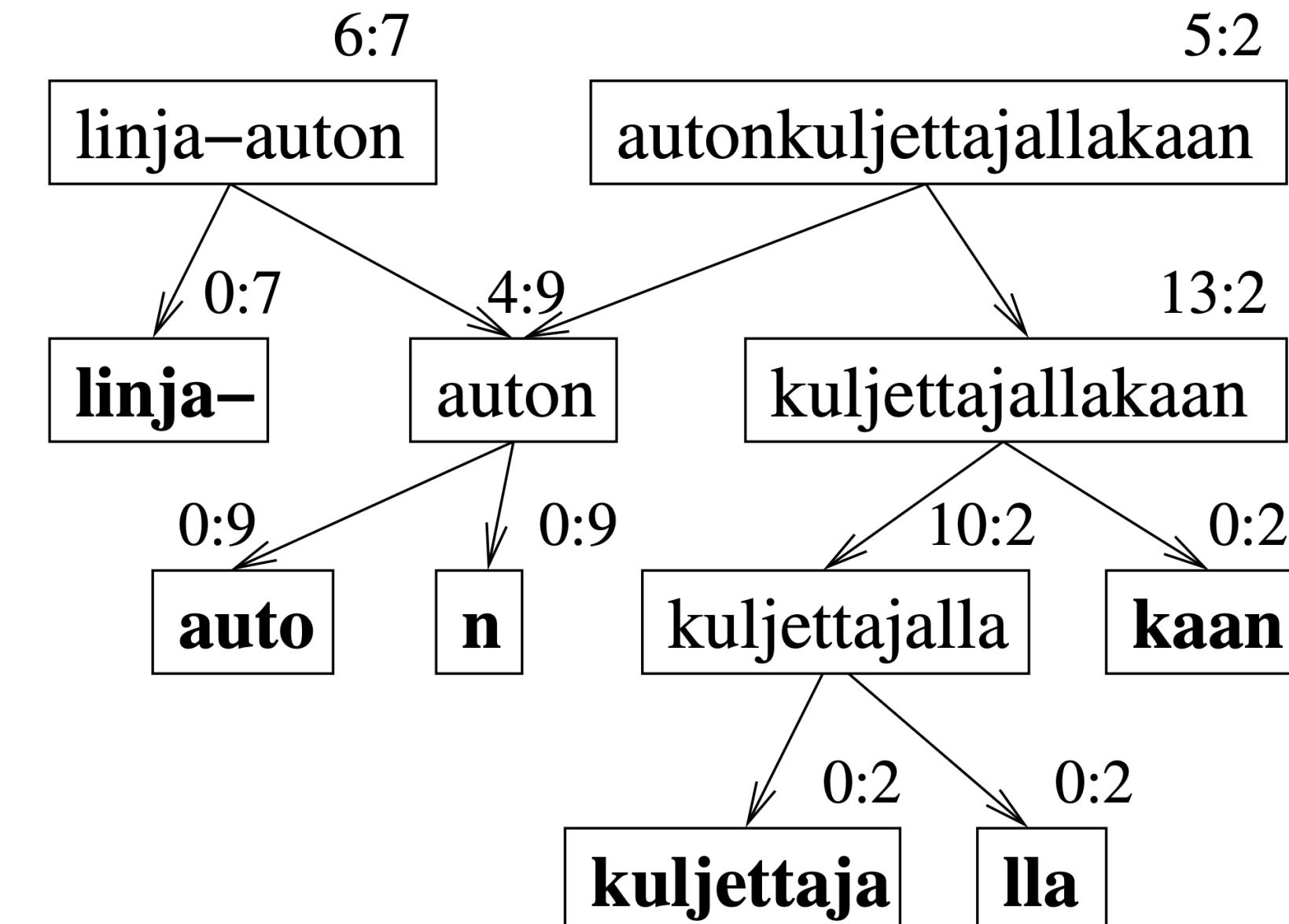
$$\begin{aligned} C &= \text{Cost(Source text)} + \text{Cost(Codebook)} \\ &= \sum_{\text{tokens}} -\log p(m_i) + \sum_{\text{types}} k * l(m_j) \end{aligned}$$



# Sequence Segmentation

- Goal: find **meaningful sub-sequences** in language

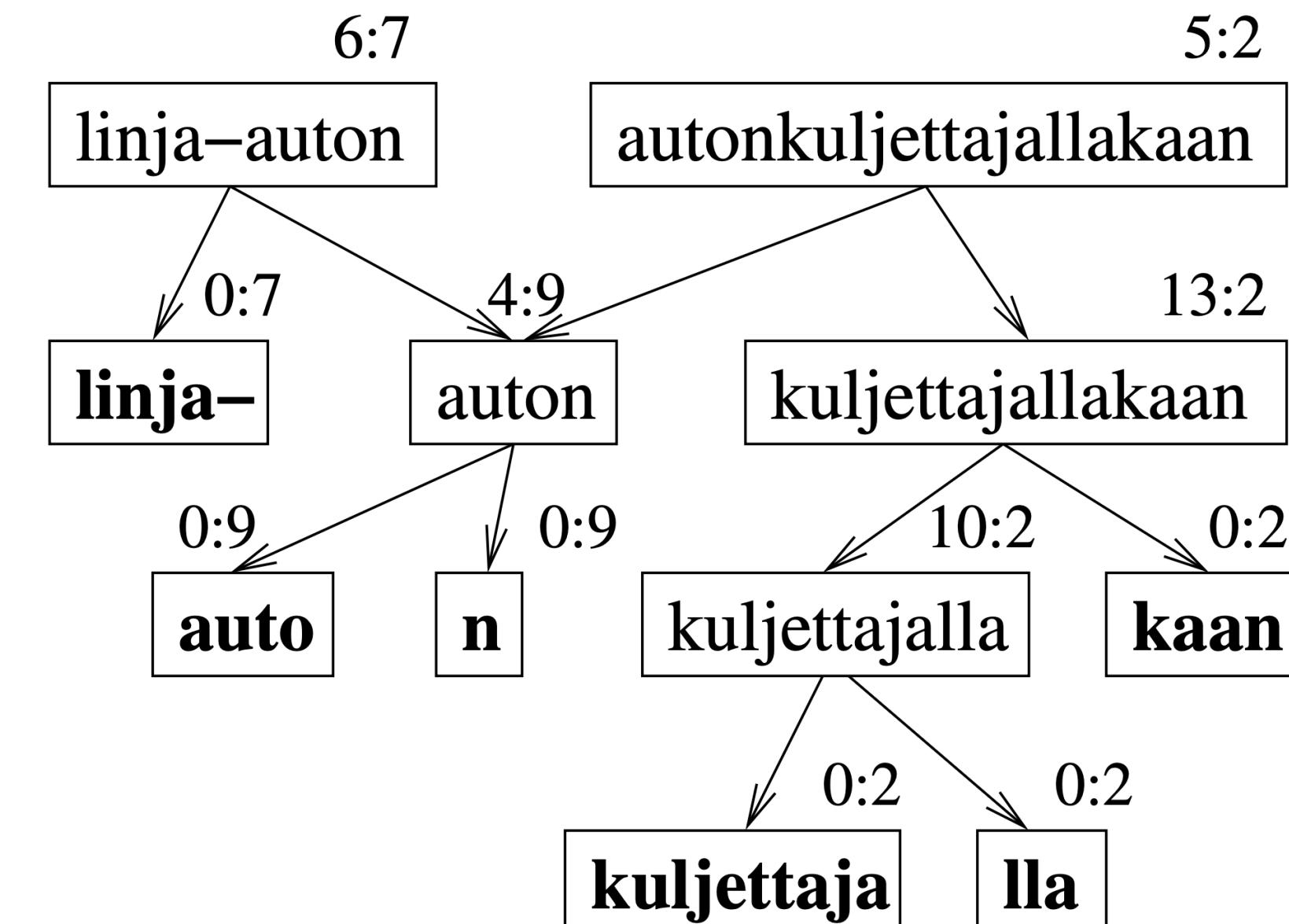
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# Sequence Segmentation

- Goal: find **meaningful sub-sequences** in language
- Surrogate metric: **Minimum Description Length (MDL)**
  - Simultaneously minimize the **data complexity** and **description length**
  - More unique, long units → **longer description** → under-segmentation

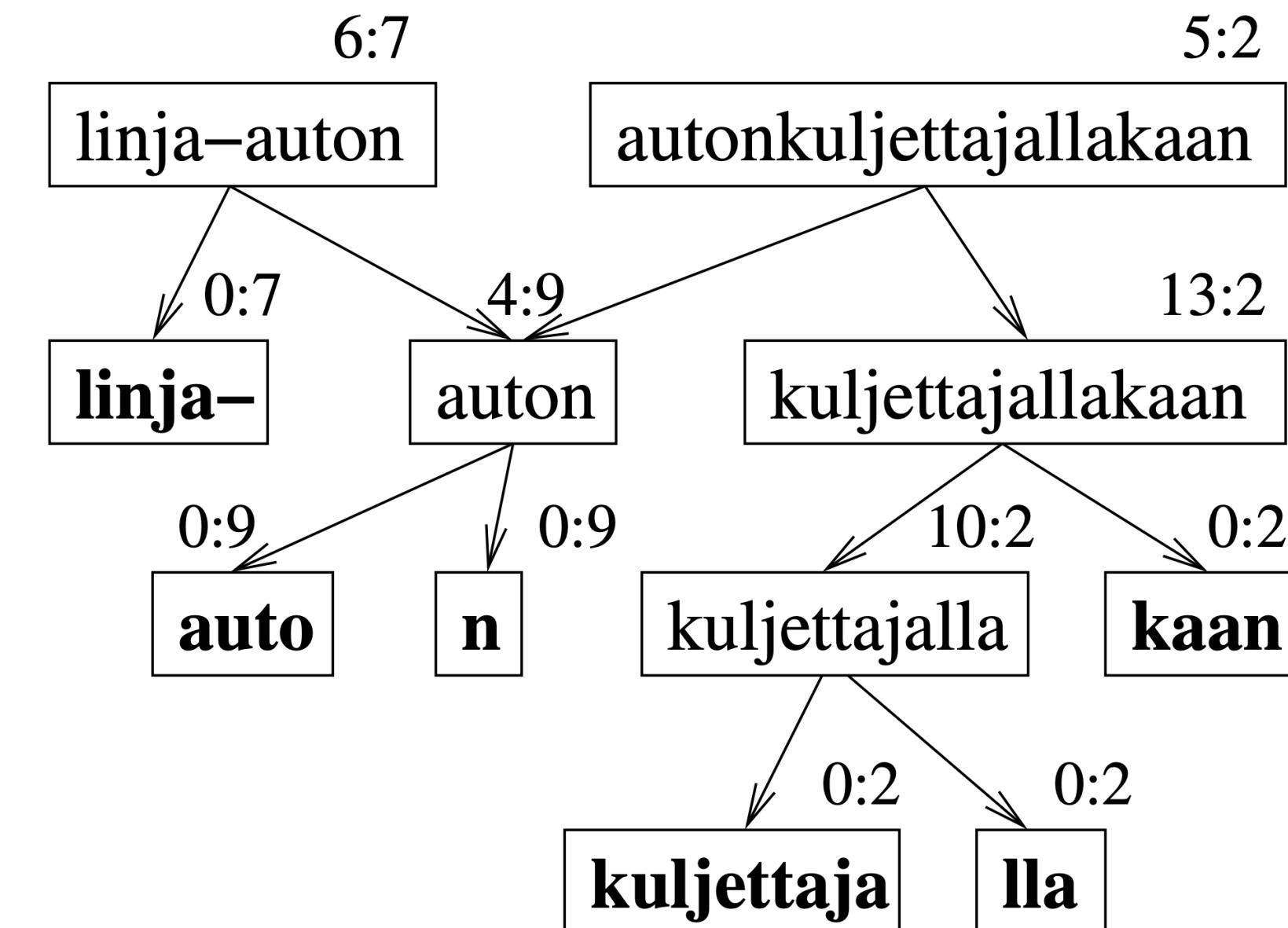
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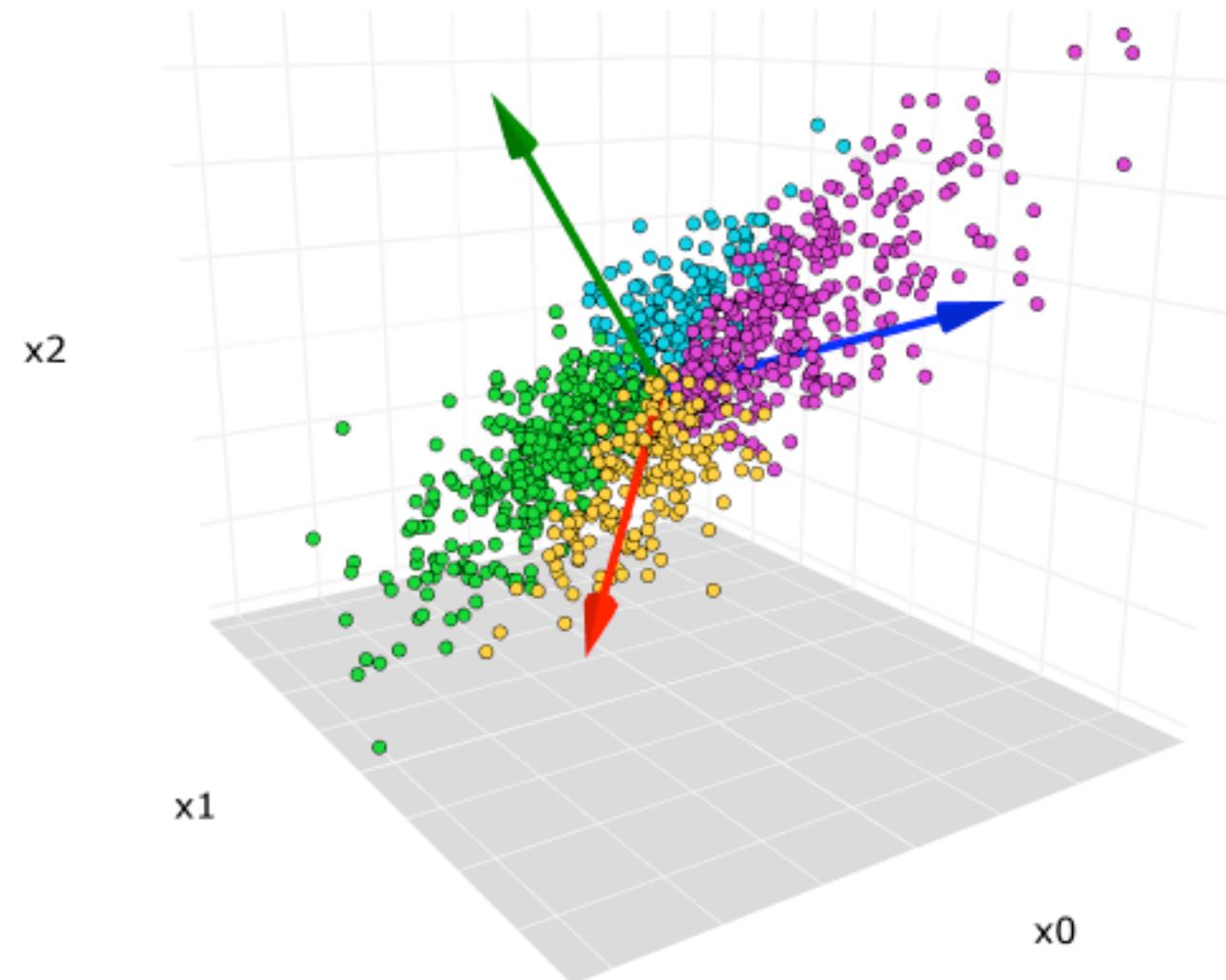
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- Inductive bias: assumes language **prefers efficiency** (not guaranteed)

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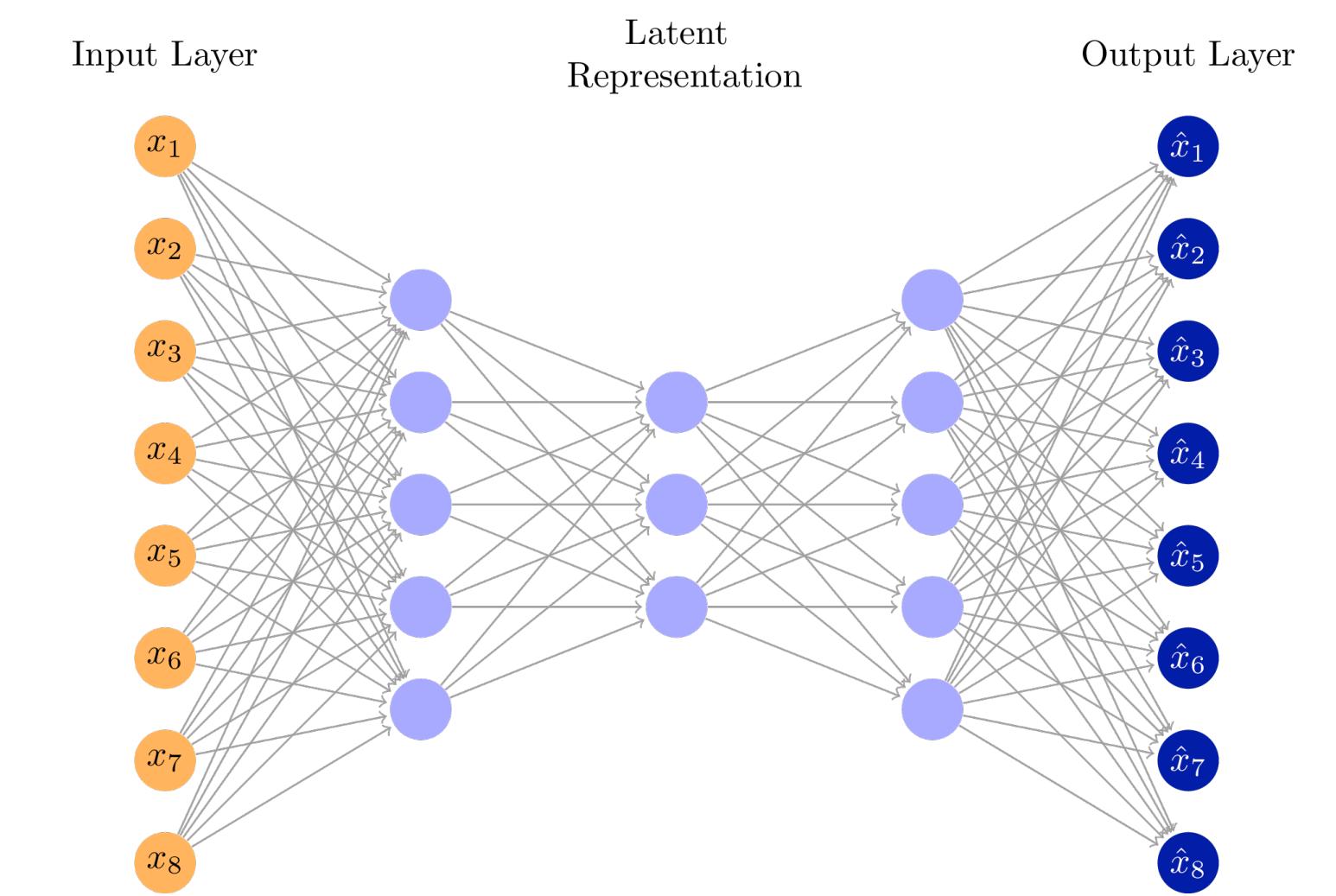
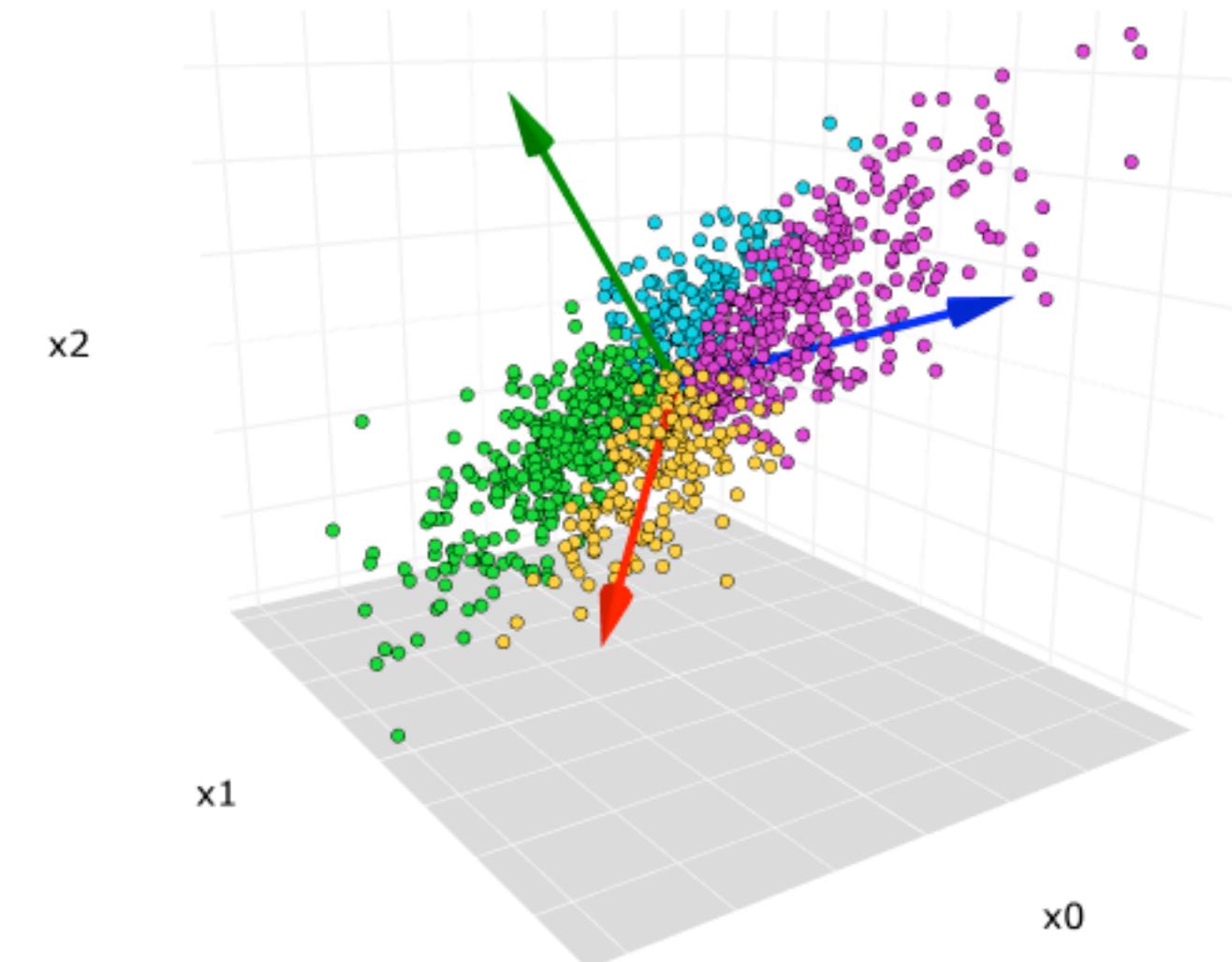


# Today: Continuous Structure



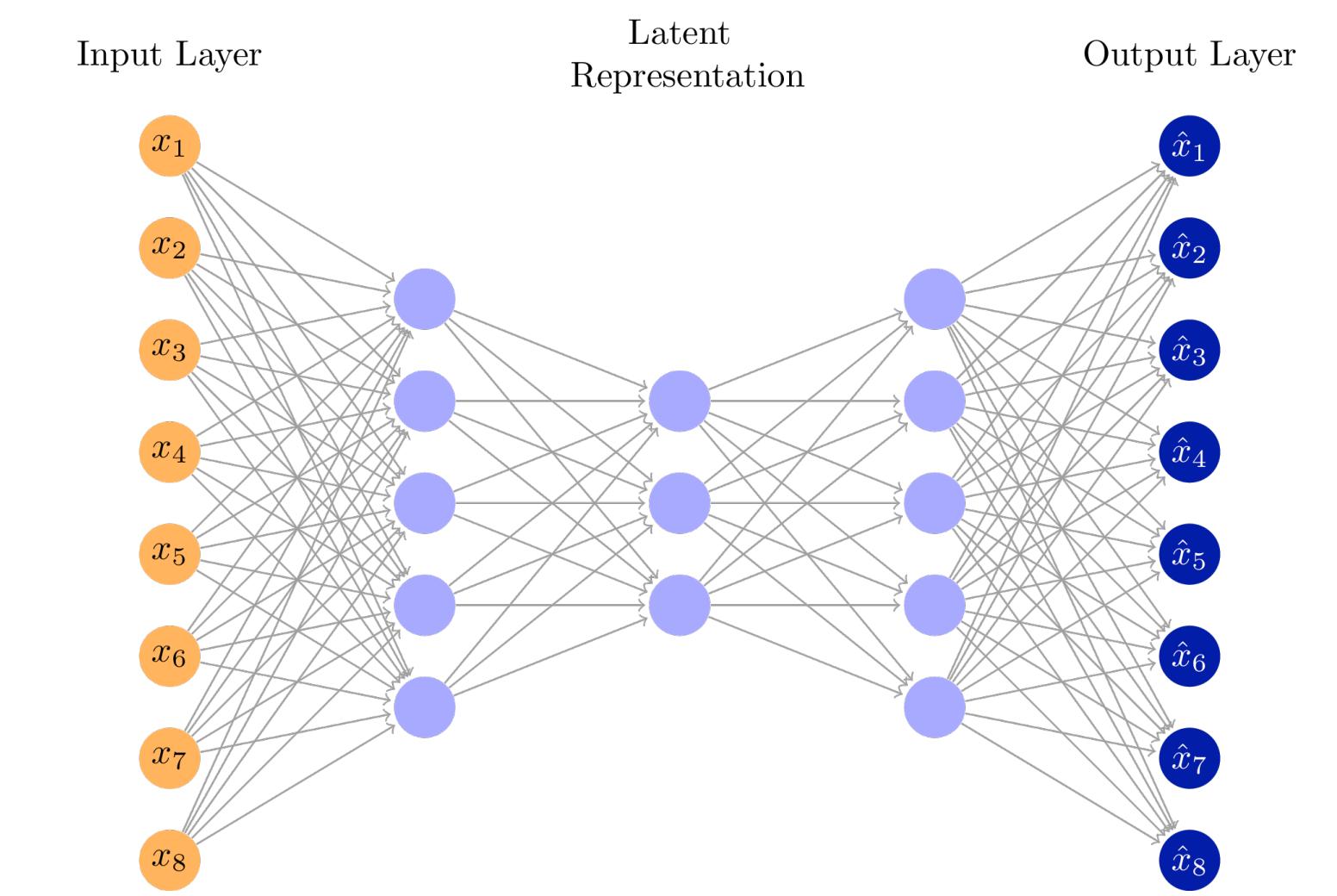
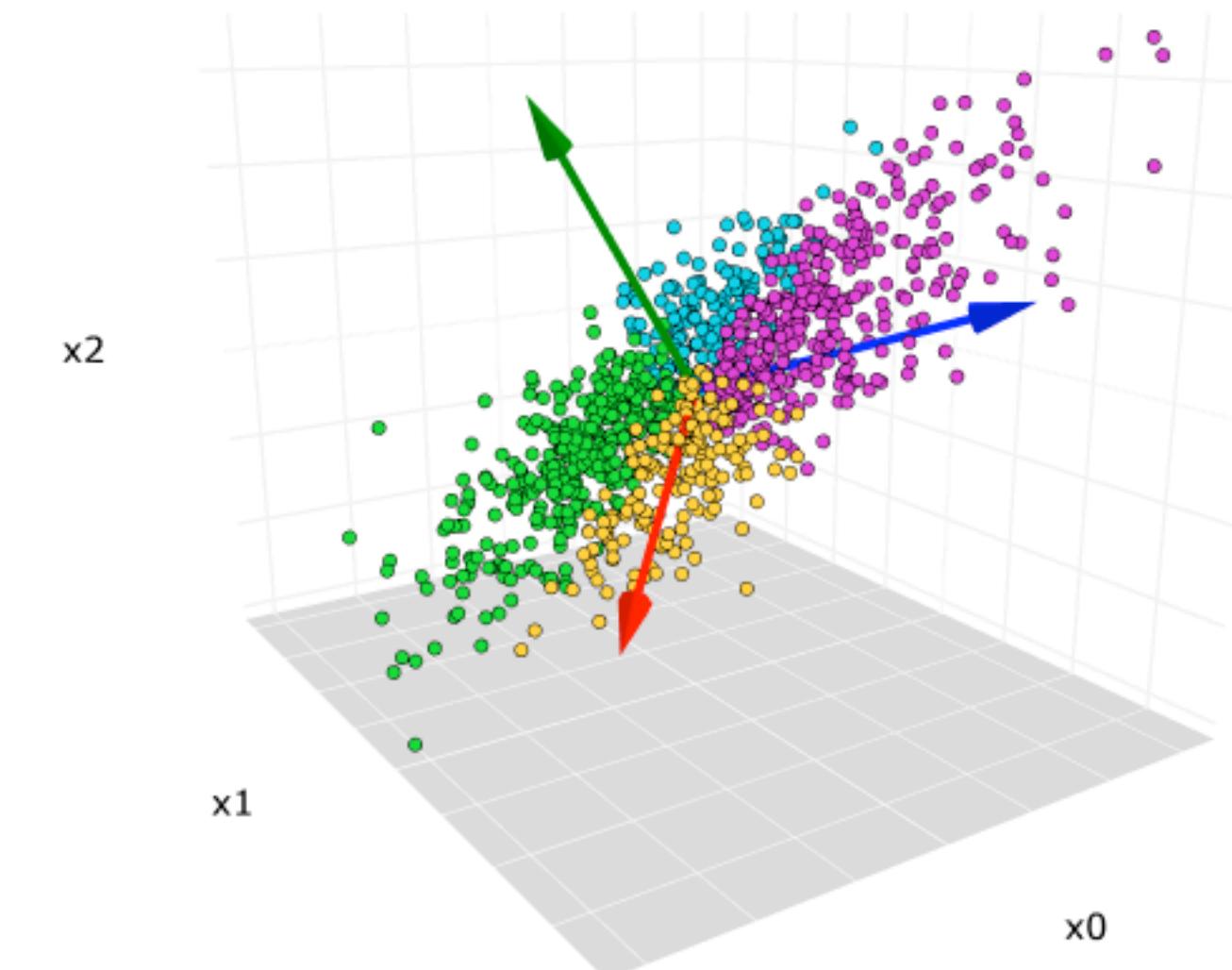
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- Instead of categorizing data, find **useful transformations**



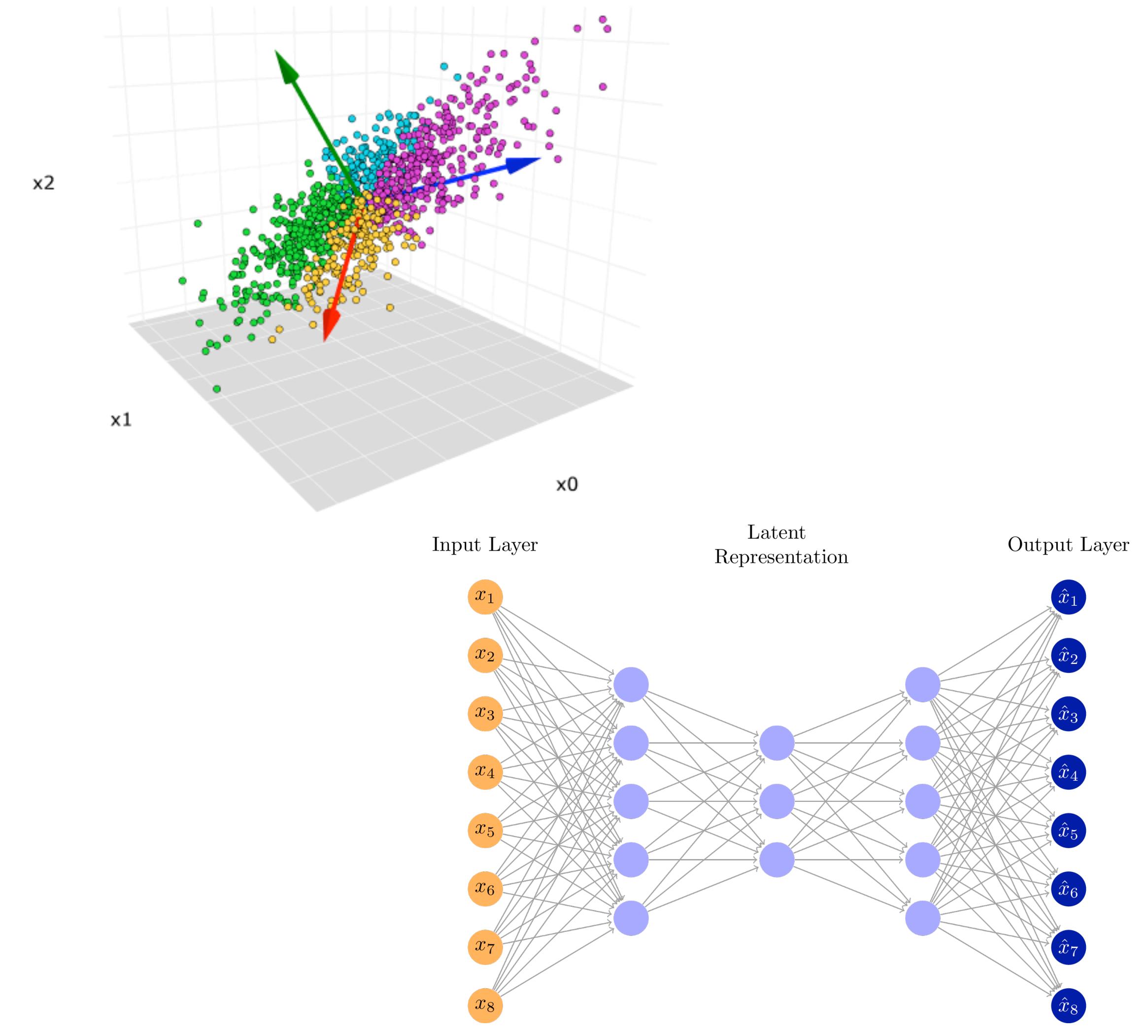
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- Instead of categorizing data, find **useful transformations**
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  - Either via **explaining variance** or **minimizing reconstruction error**



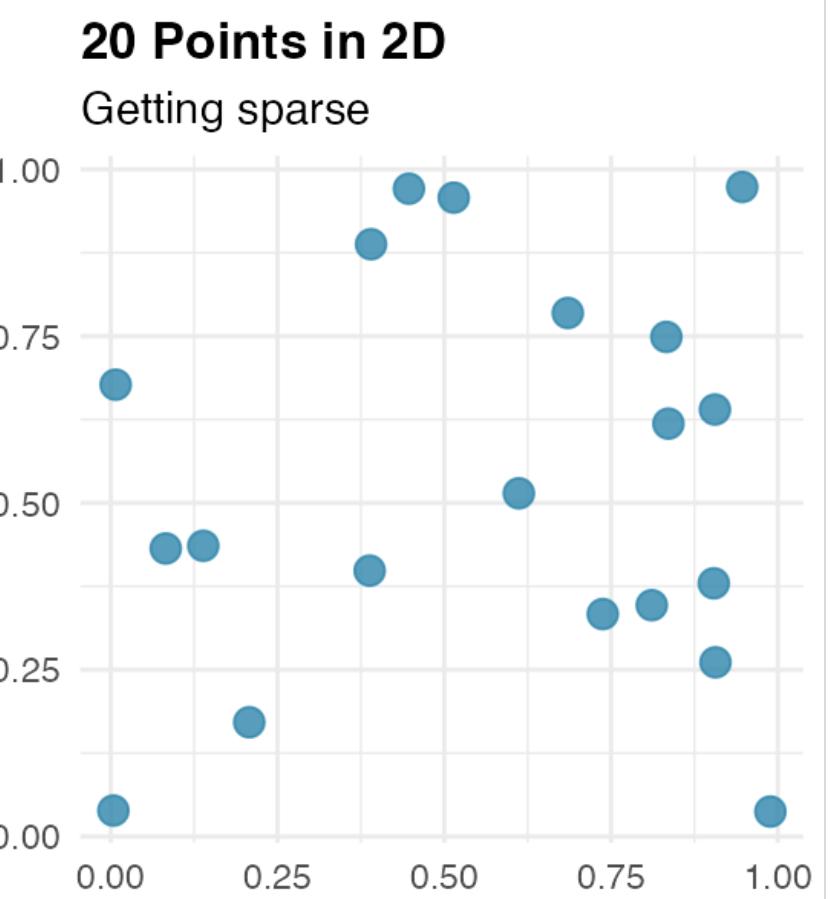
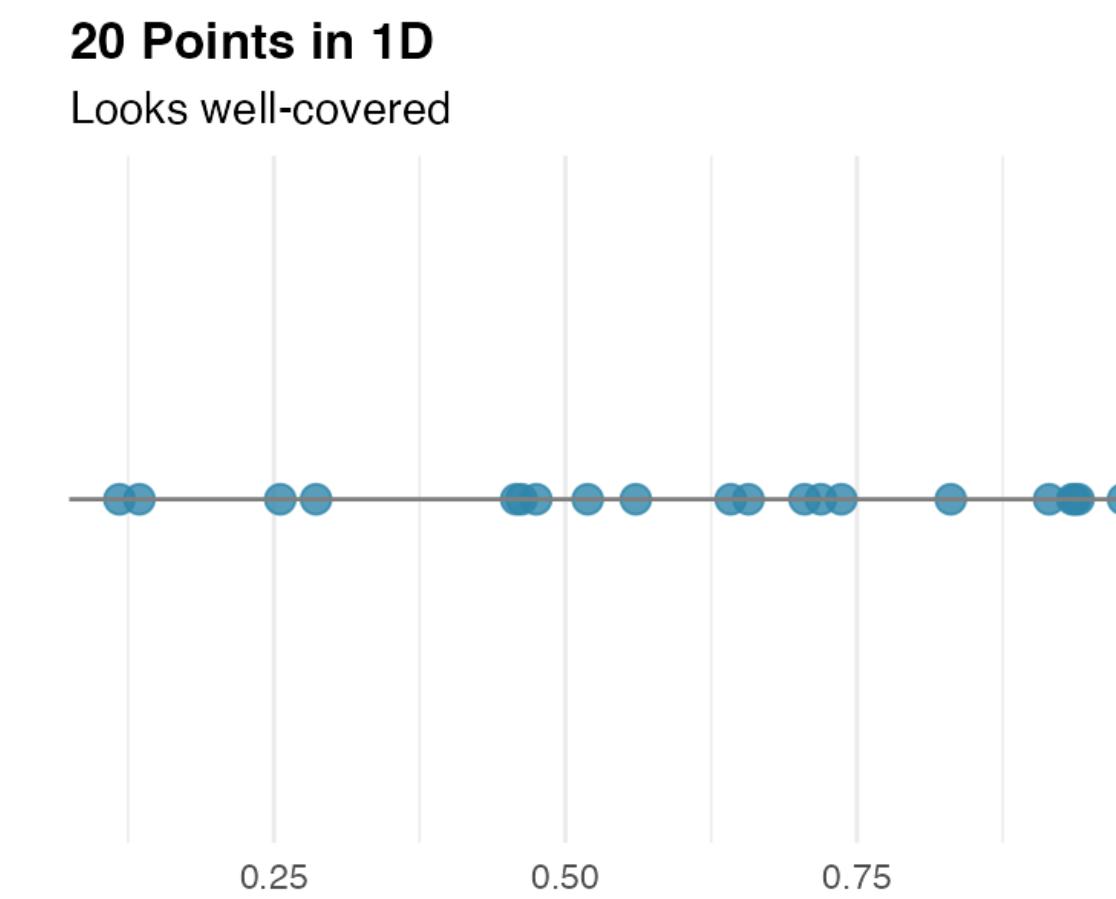
# Today: Continuous Structure

- Instead of categorizing data, find **useful transformations**
- Surrogate objectives: **dimensionality reduction**
  - Either via **explaining variance** or **minimizing reconstruction error**
- The hope: remaining dims are **more useful for downstream tasks**
  - Inductive bias that **most dimensions are redundant/noise**

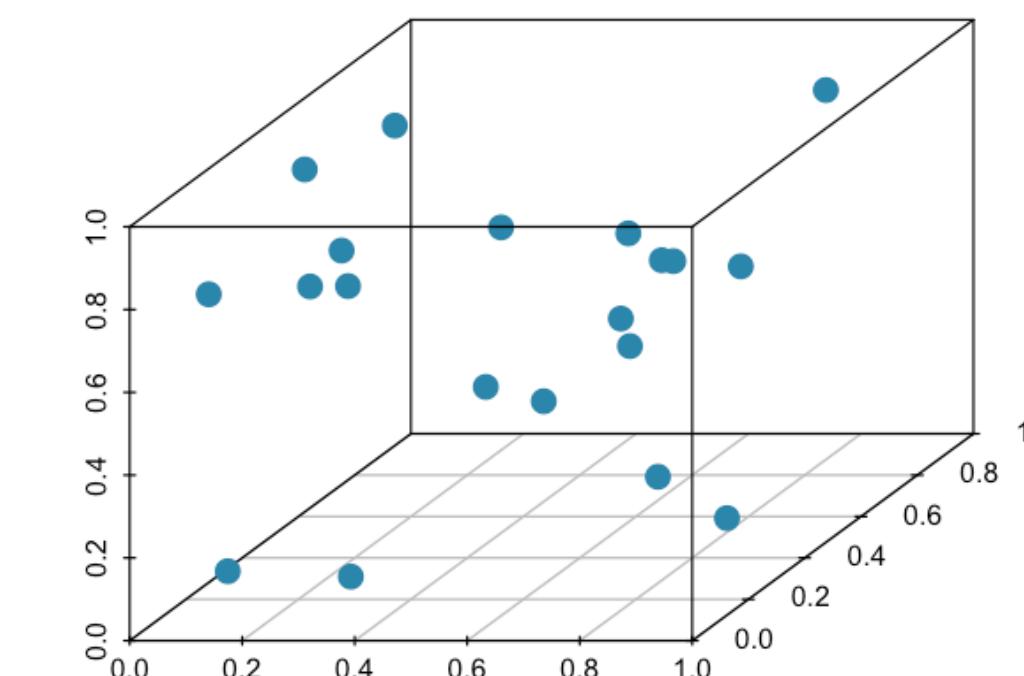


# Dimensionality Reduction

# The "Curse of Dimensionality"



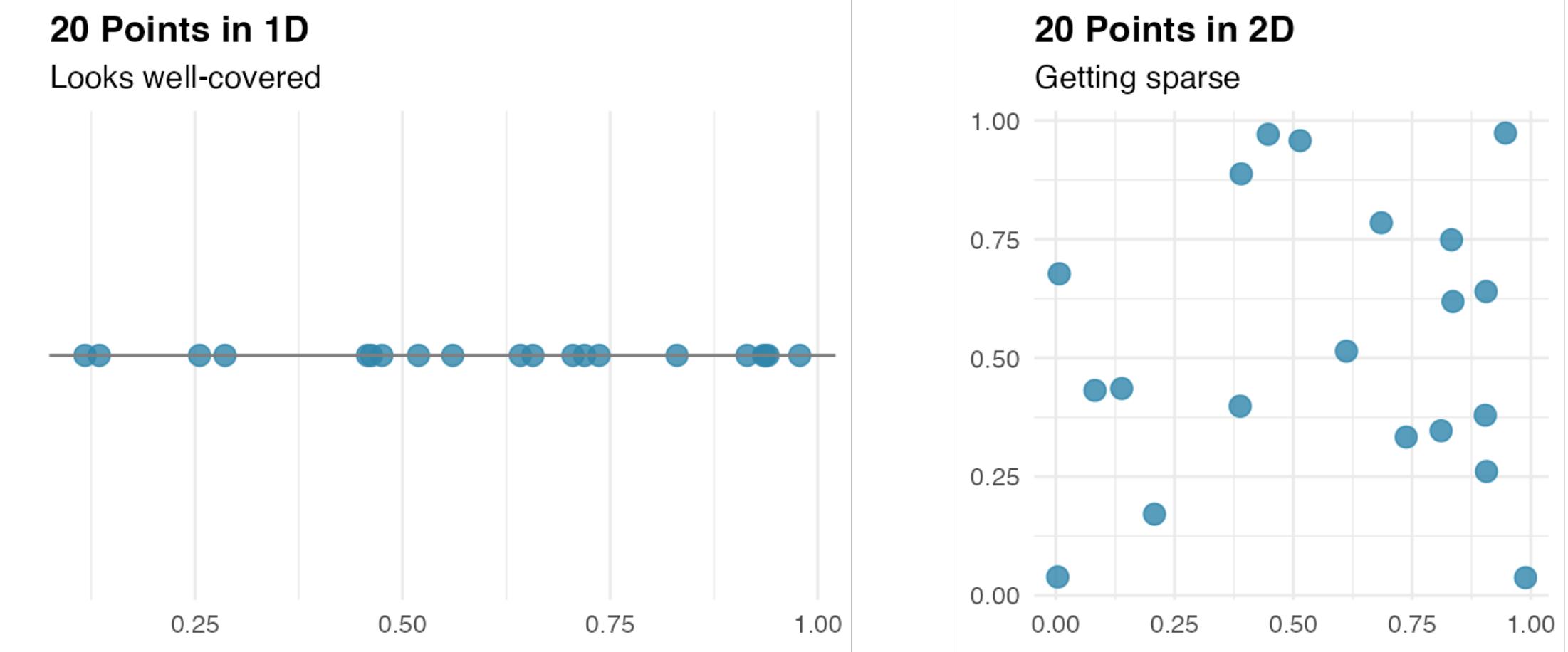
**20 Points in 3D: Very Sparse**



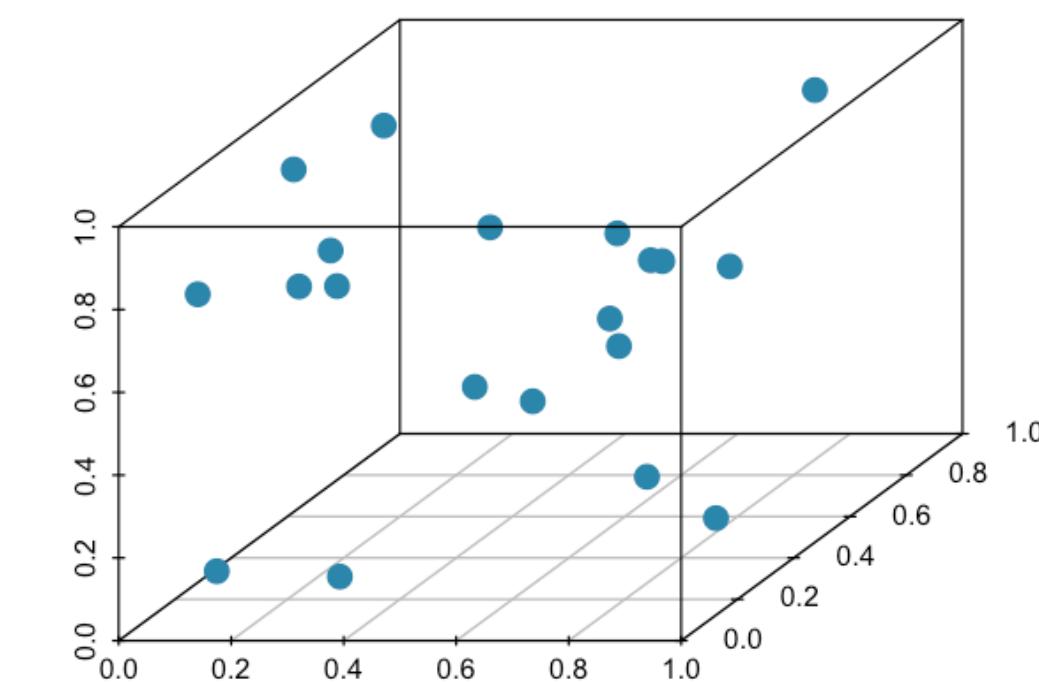
Most of the cube is empty. Now imagine 100D...

# The "Curse of Dimensionality"

- High-dimensional data is very **sparse** (i.e. empty)



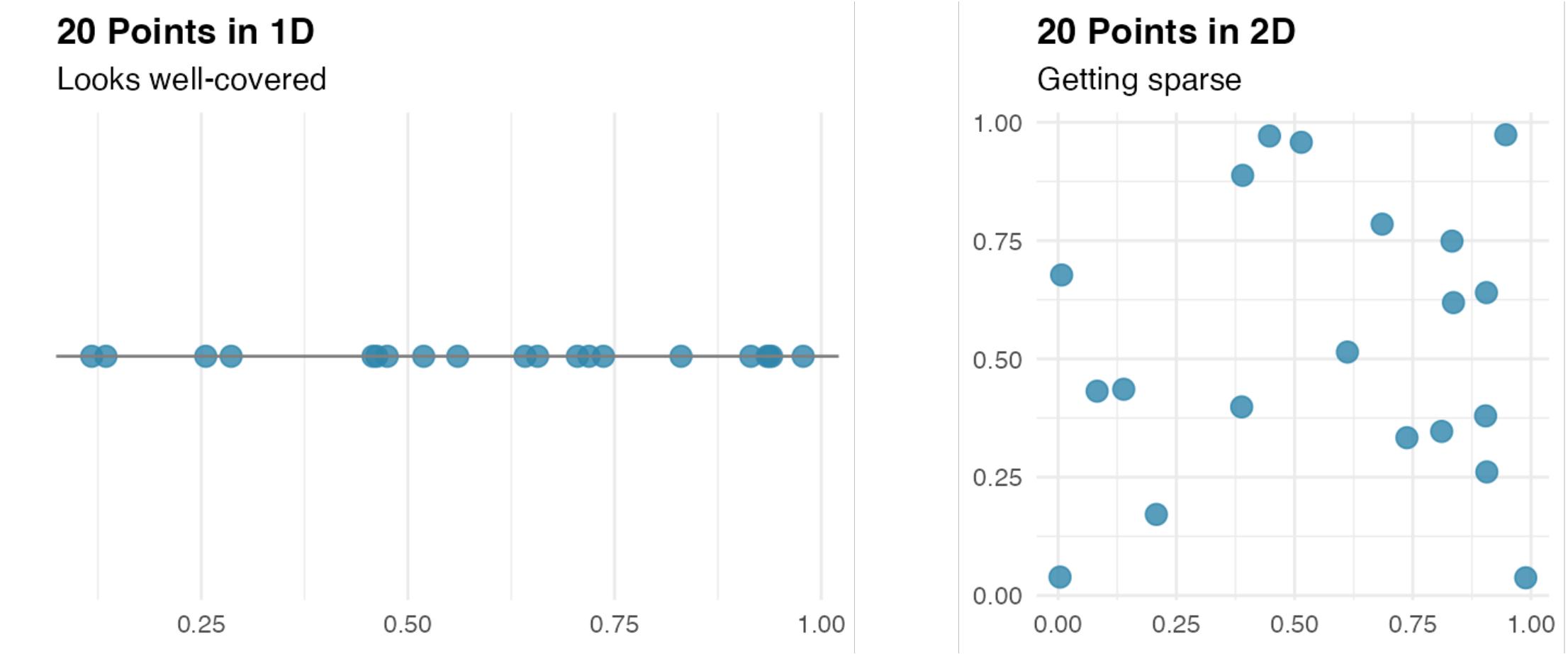
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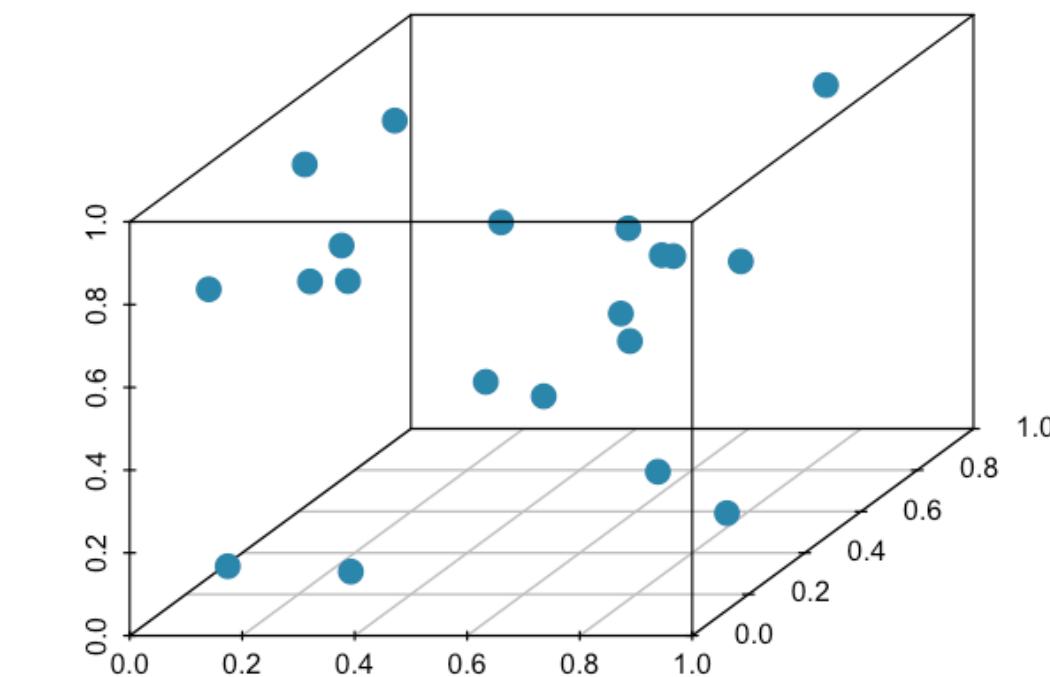
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# The "Curse of Dimensionality"

- High-dimensional data is very **sparse** (i.e. empty)
  - Say you have **50 datapoints**, spread across...



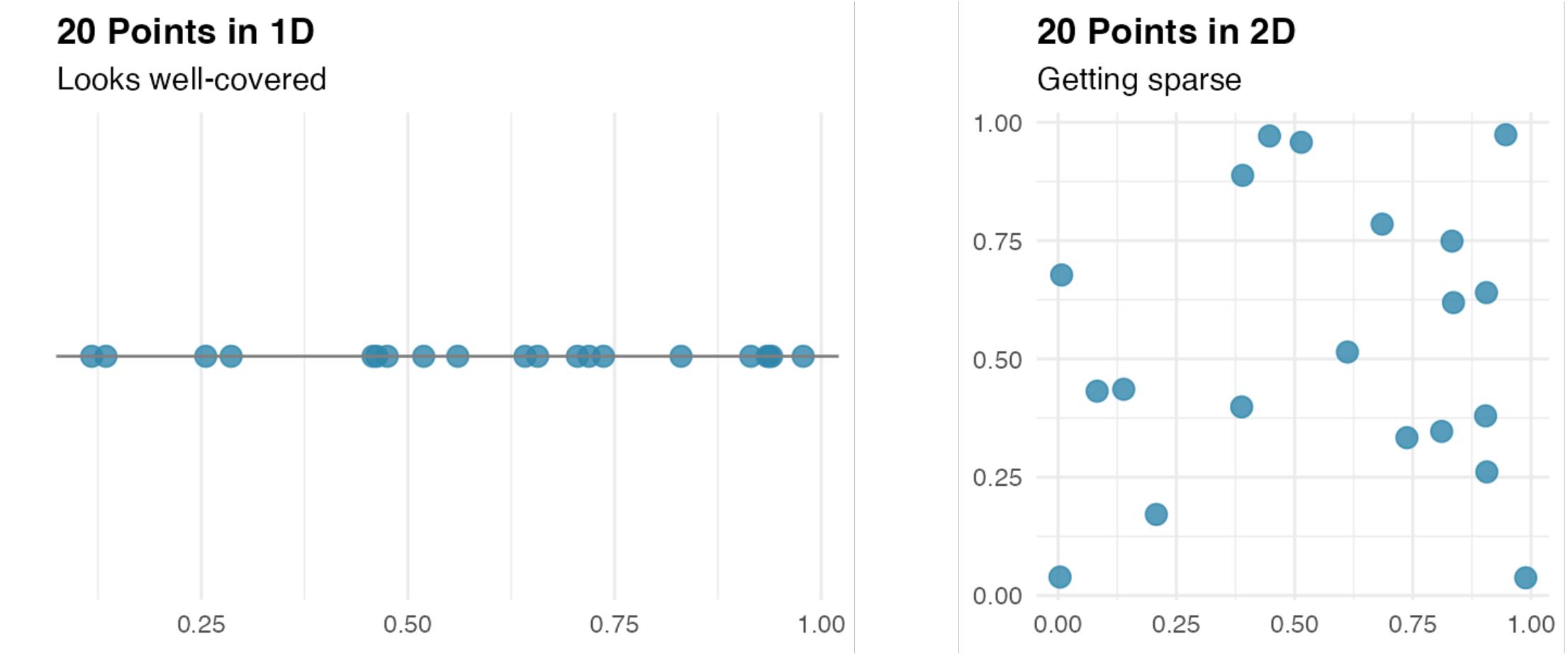
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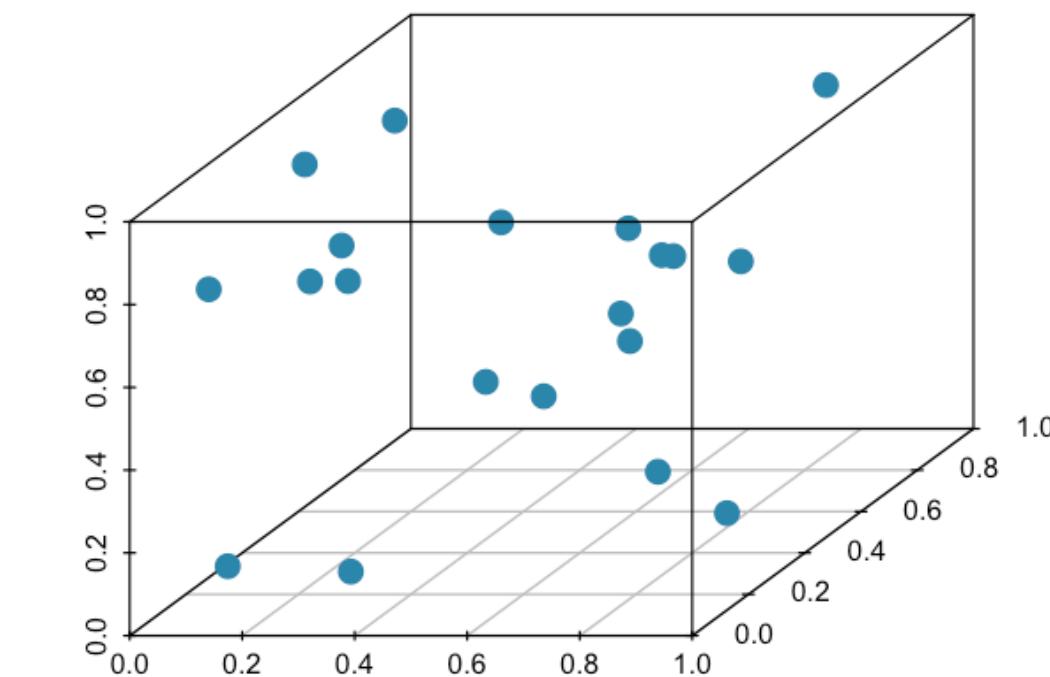
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  - A **10x10 grid** → half full



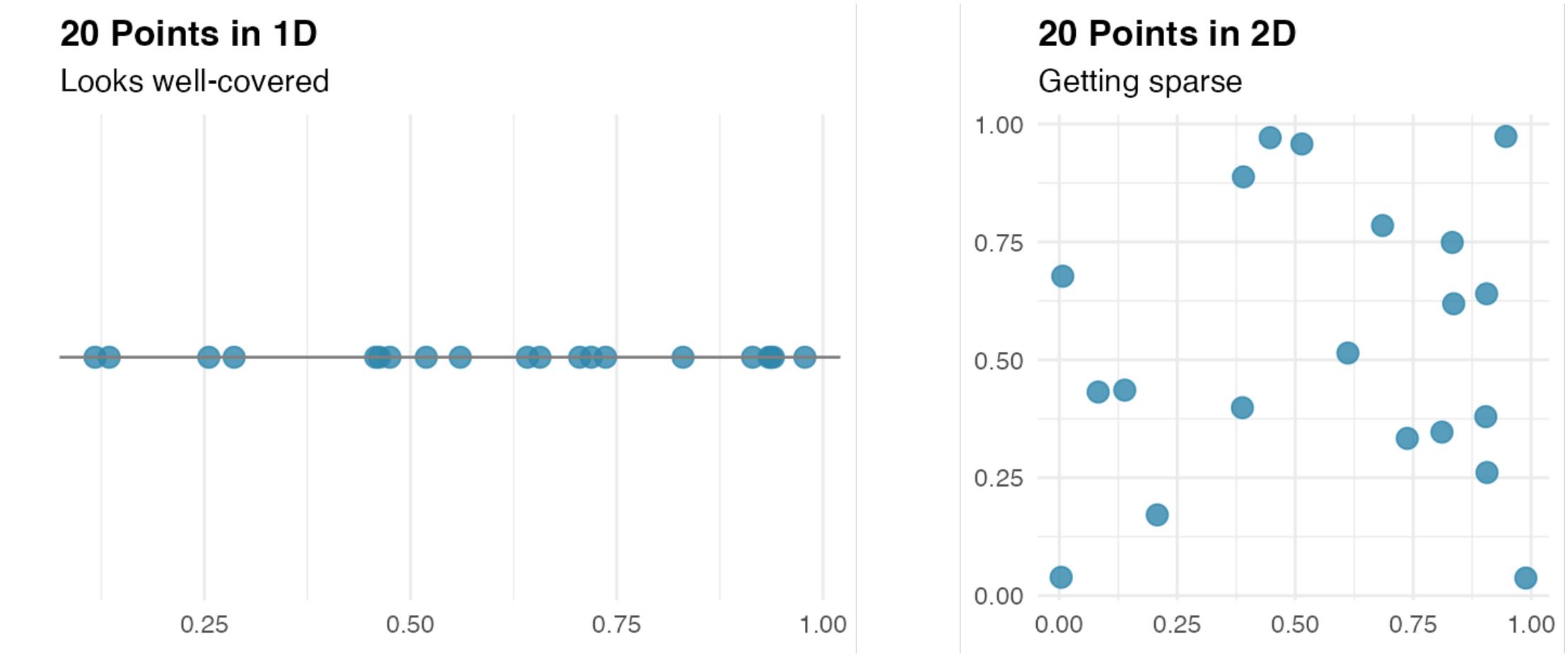
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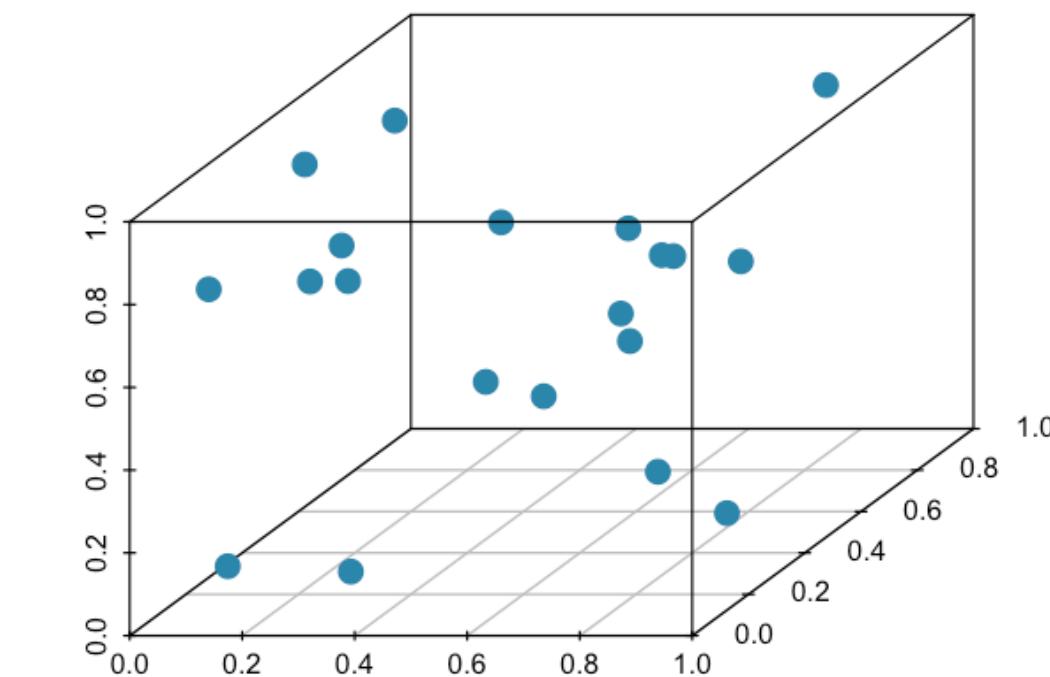
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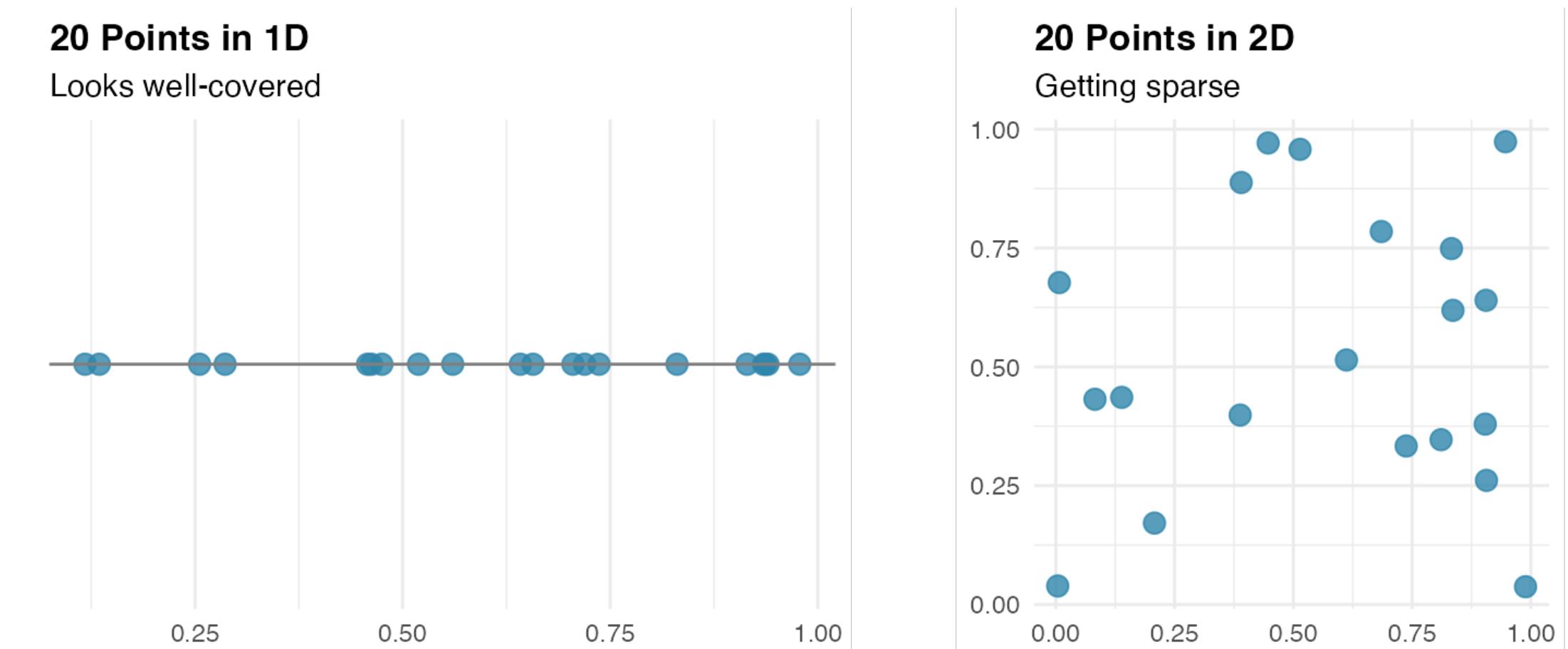
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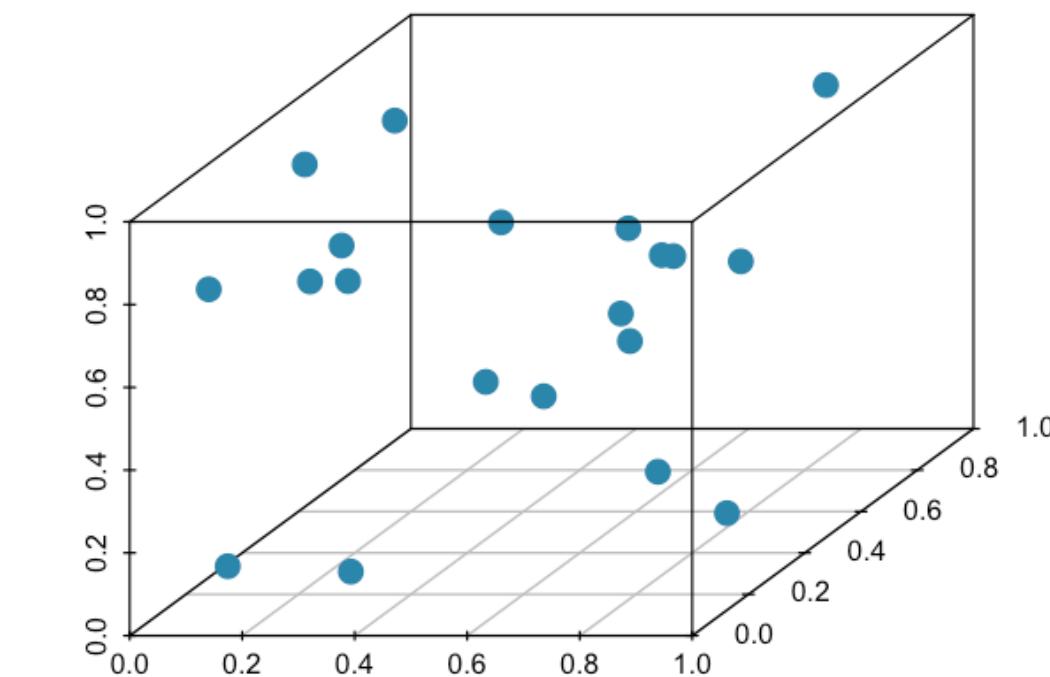
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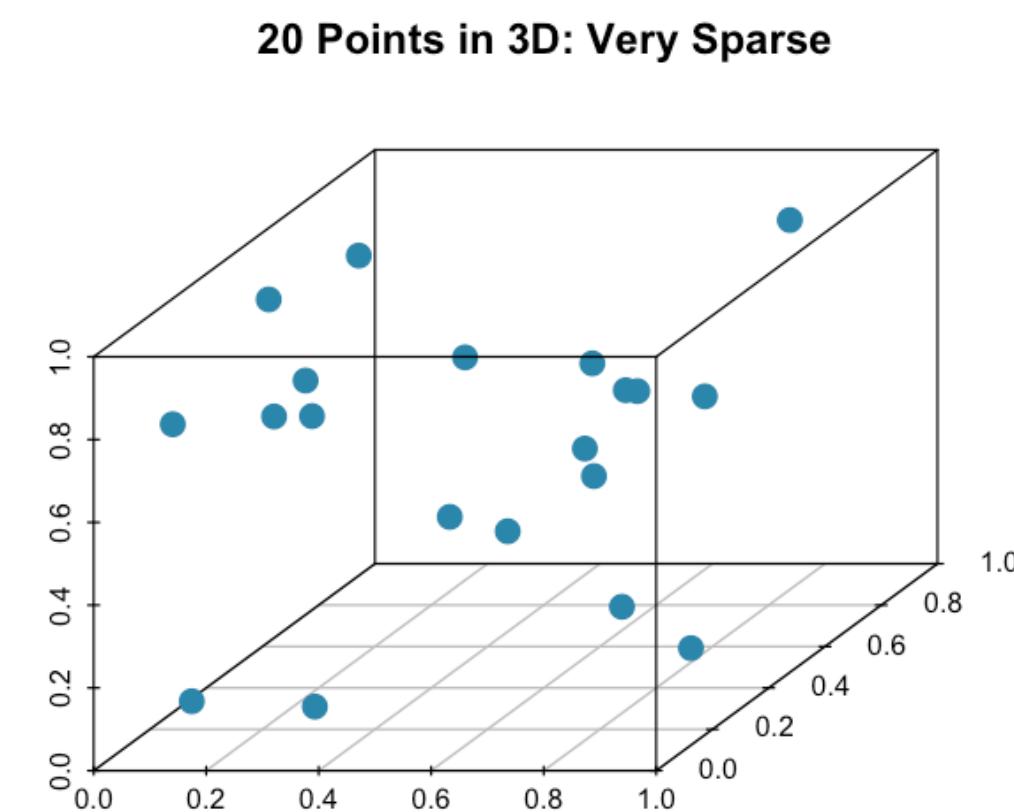
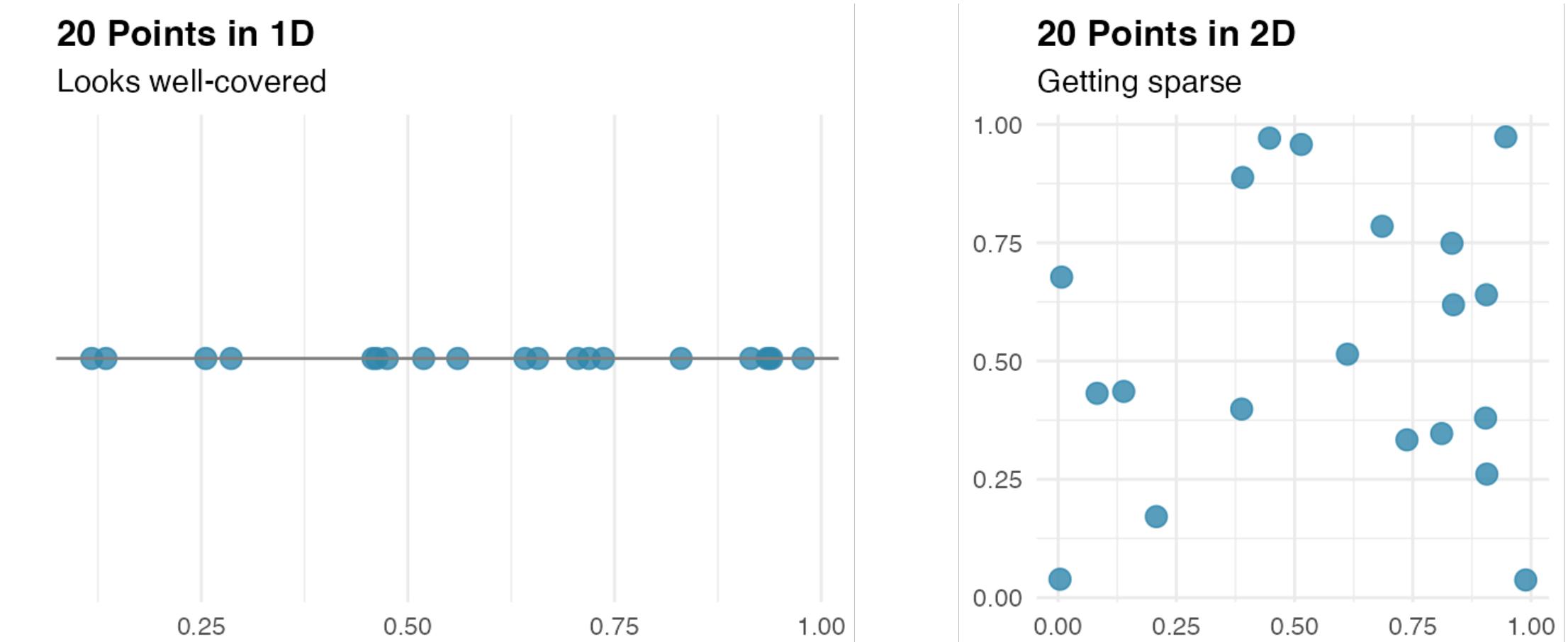
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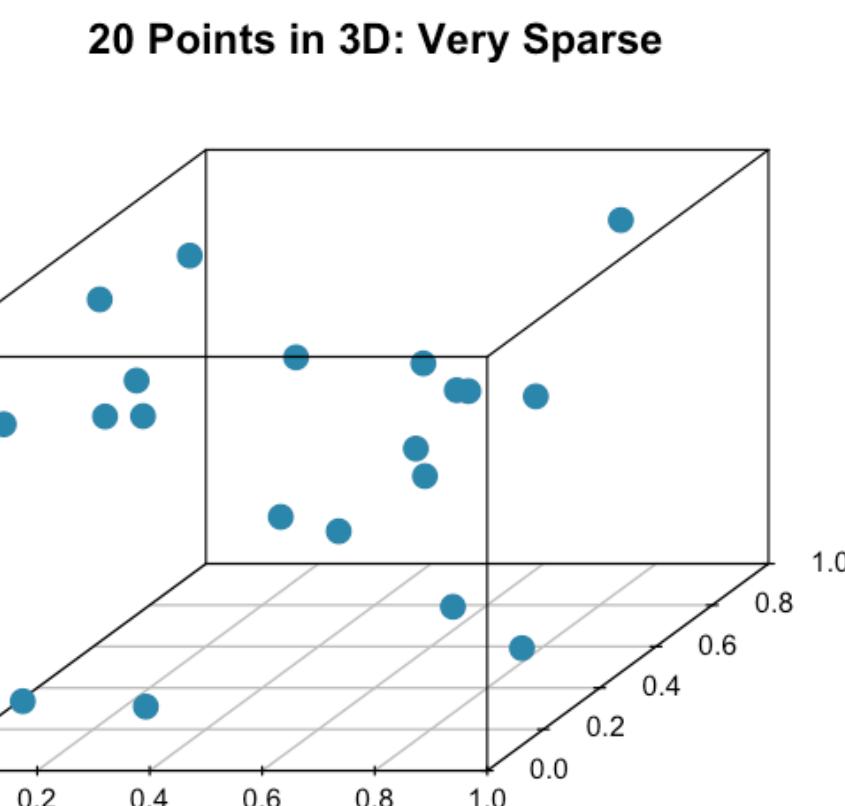
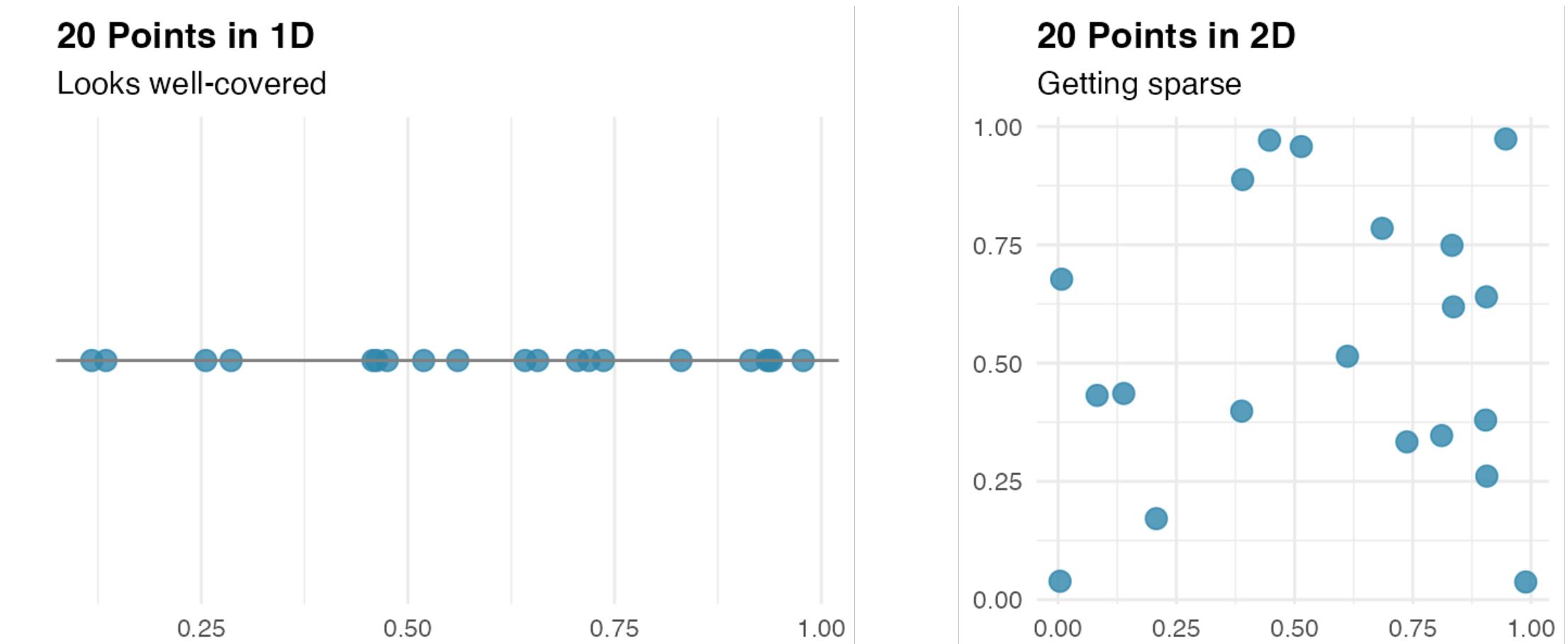
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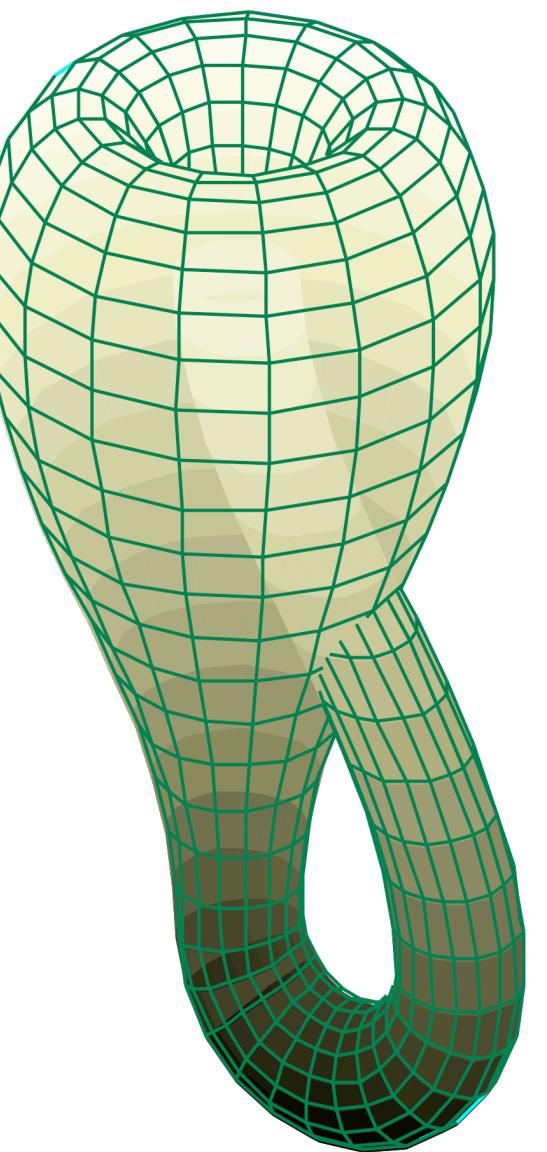
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- Calculating **distances** between points becomes **meaningless**
- Models of this space become **complex** and **data-hungry**

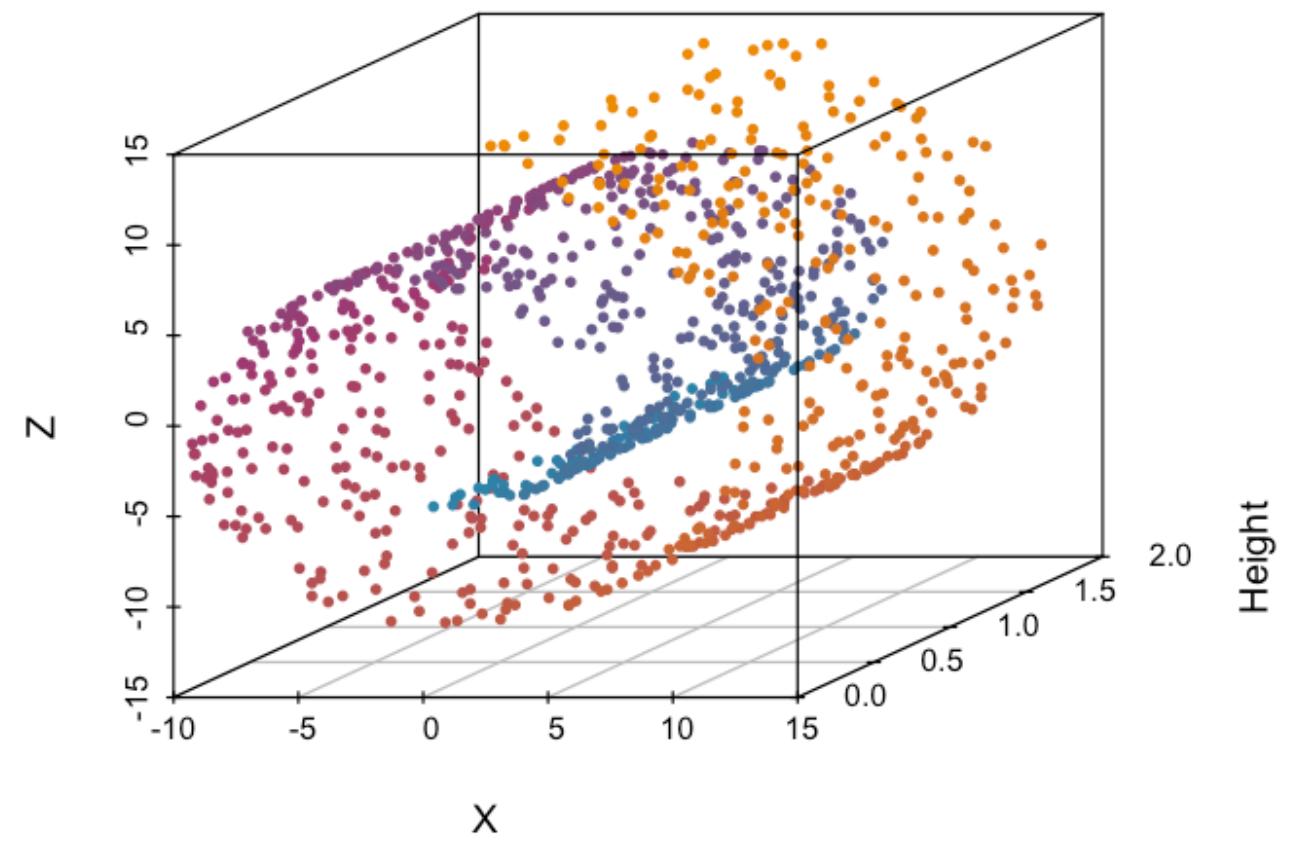


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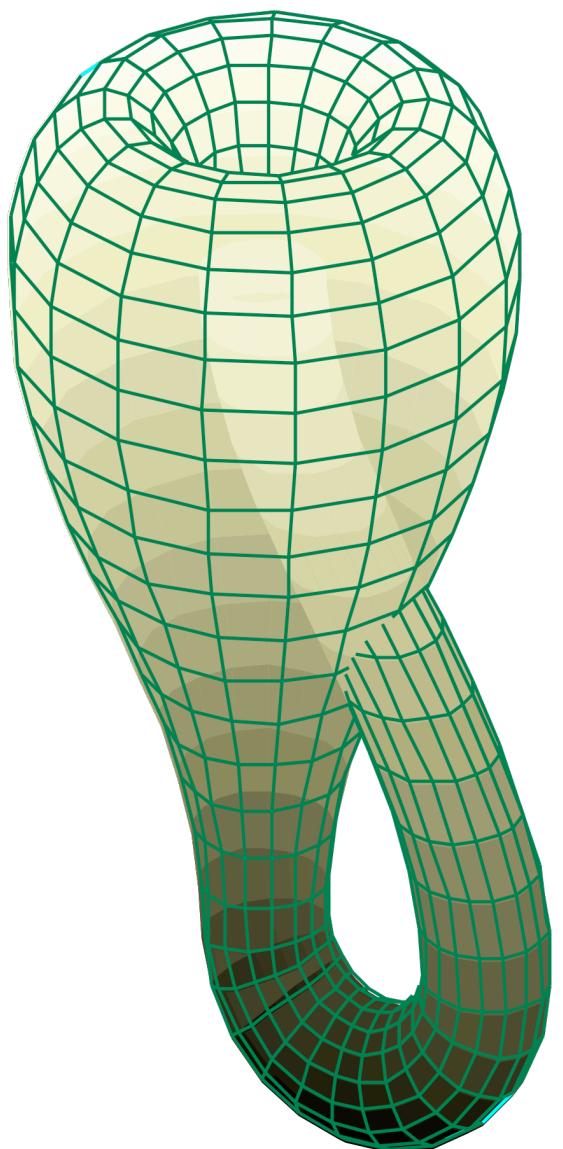


Rolled Up: A 2D Surface in 3D

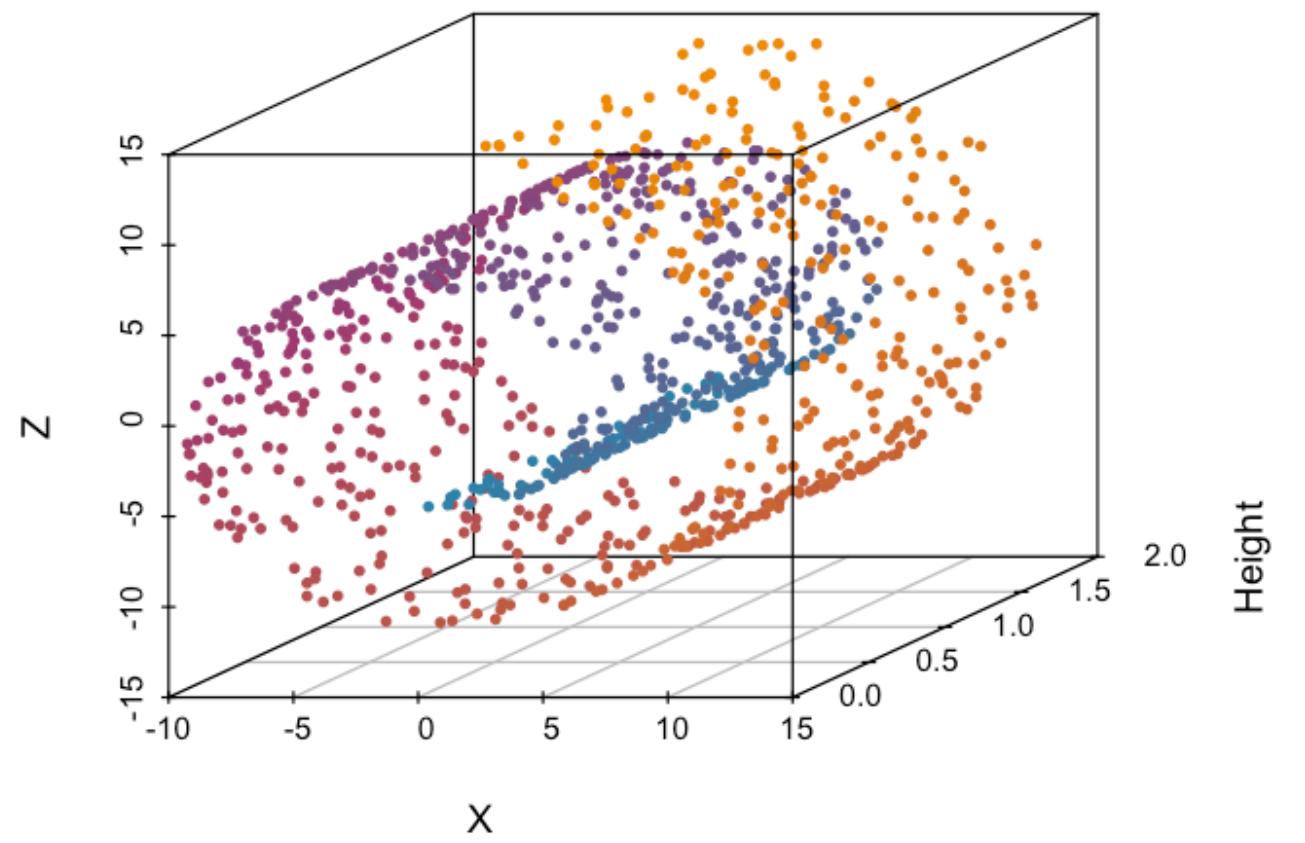


# The "Manifold Hypothesis"

- Our hope: the interesting data patterns live on a **lower-dimensional manifold** within the high-dimensional space

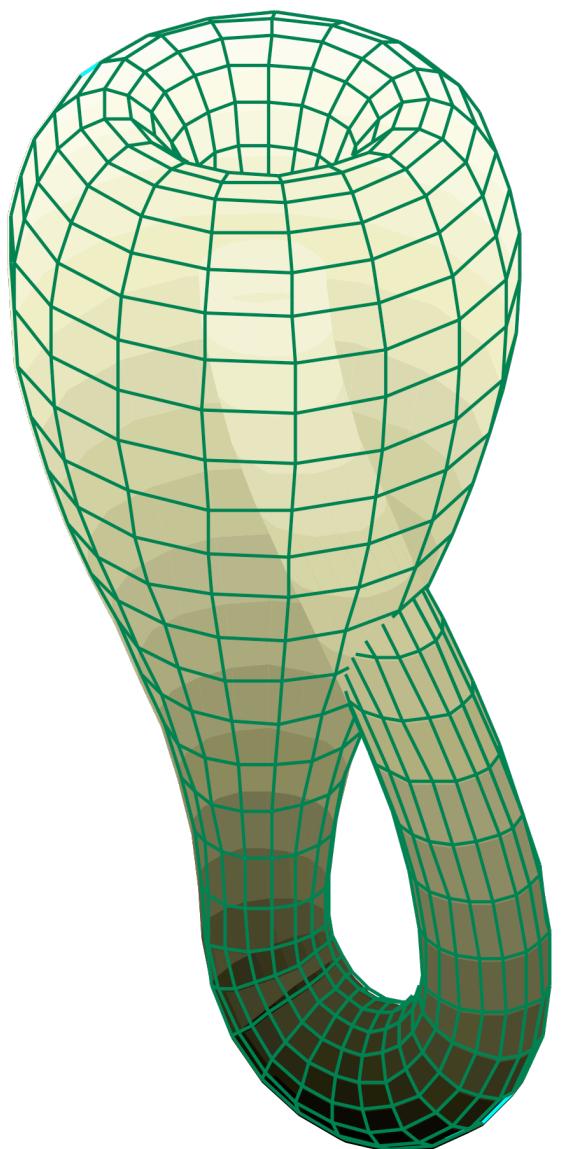


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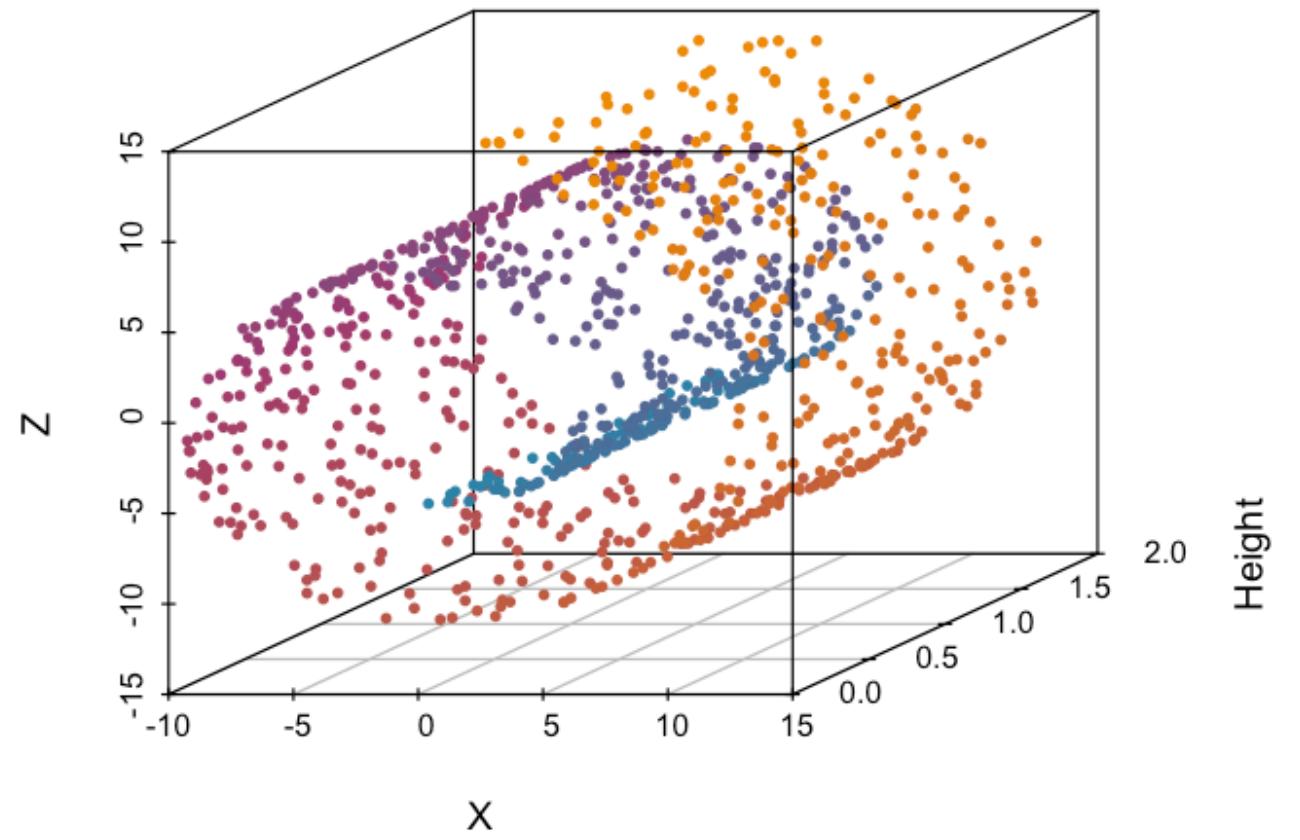


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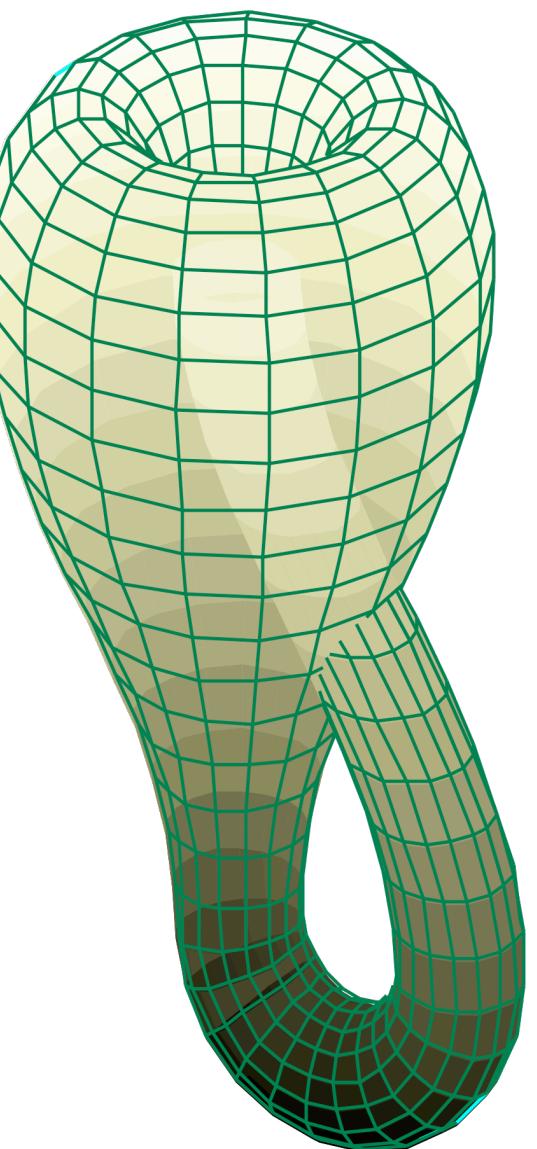


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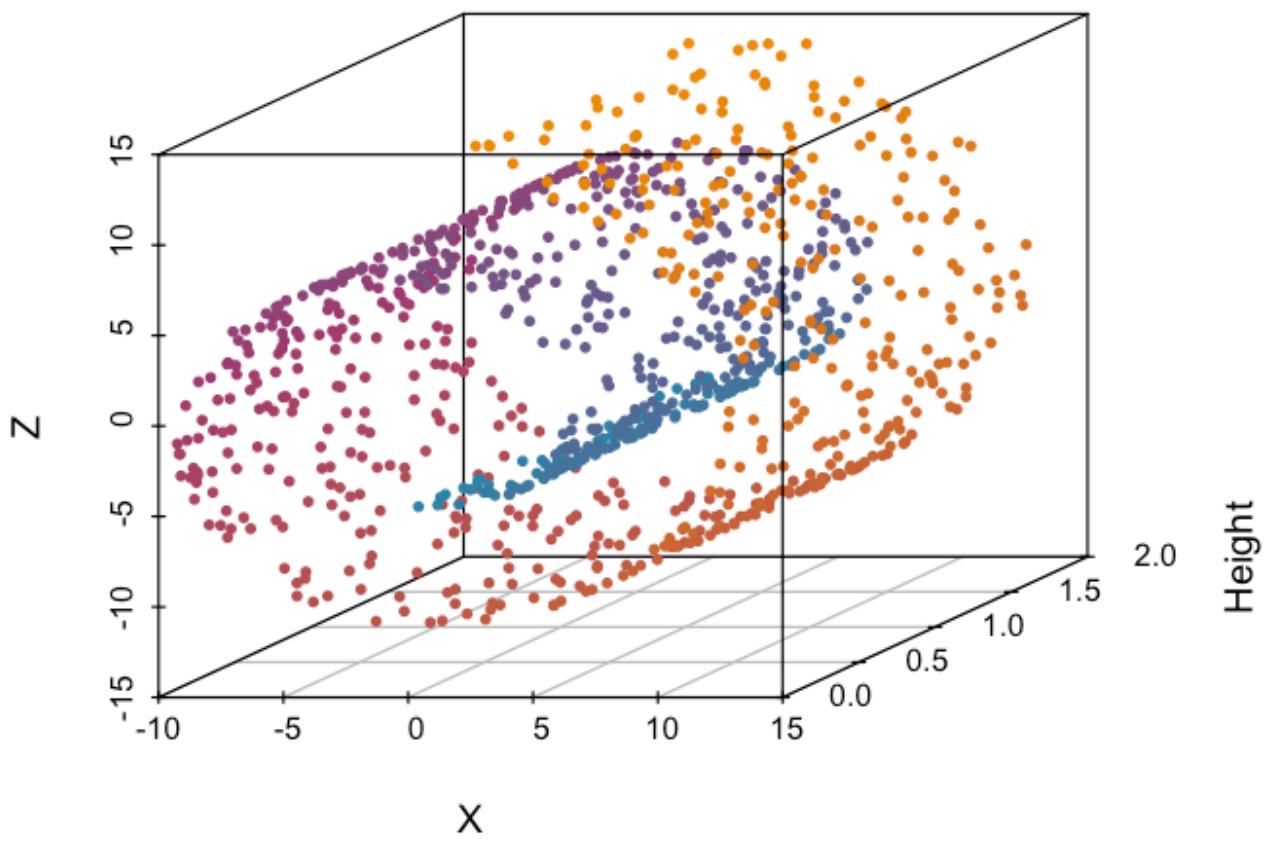


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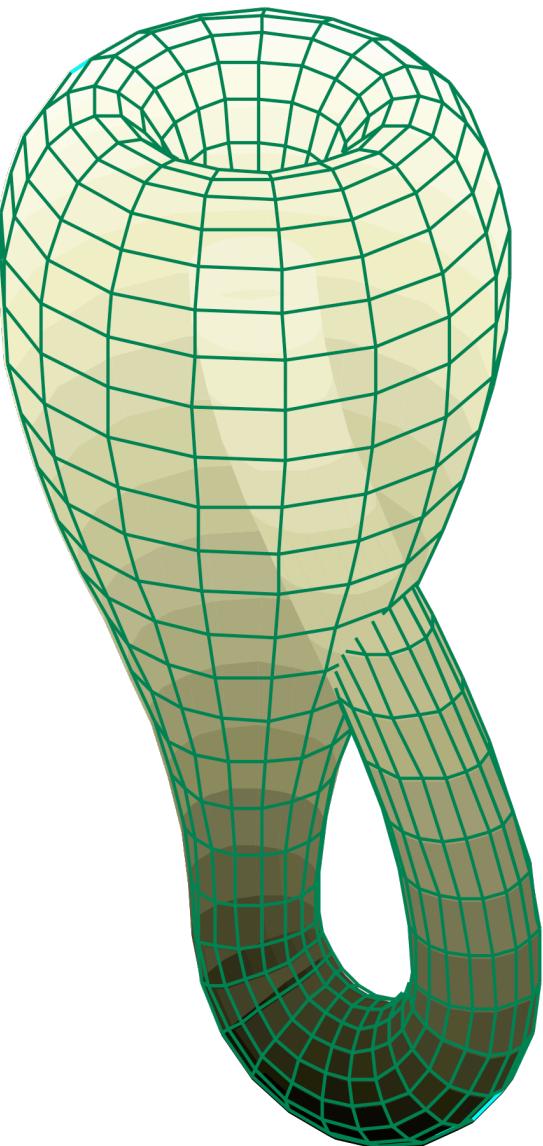


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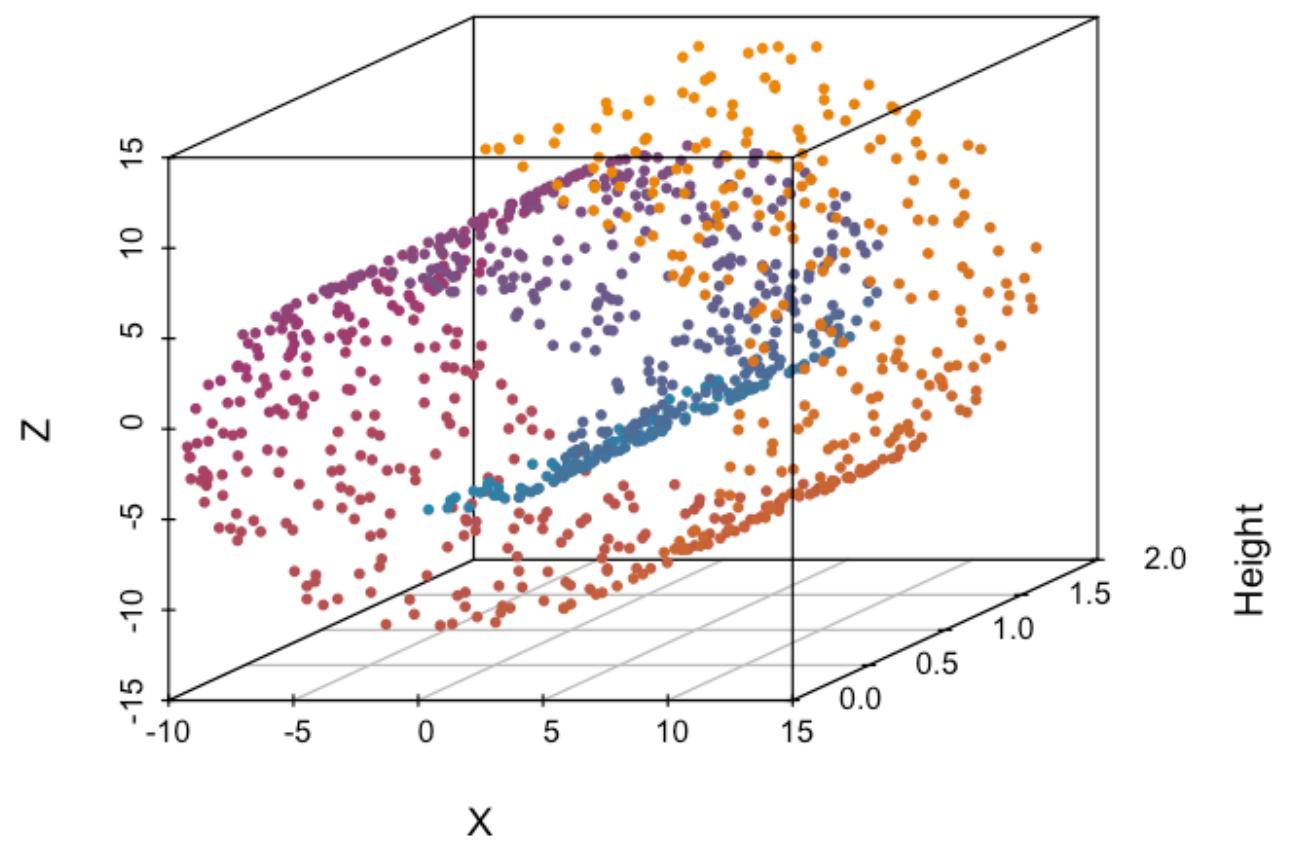


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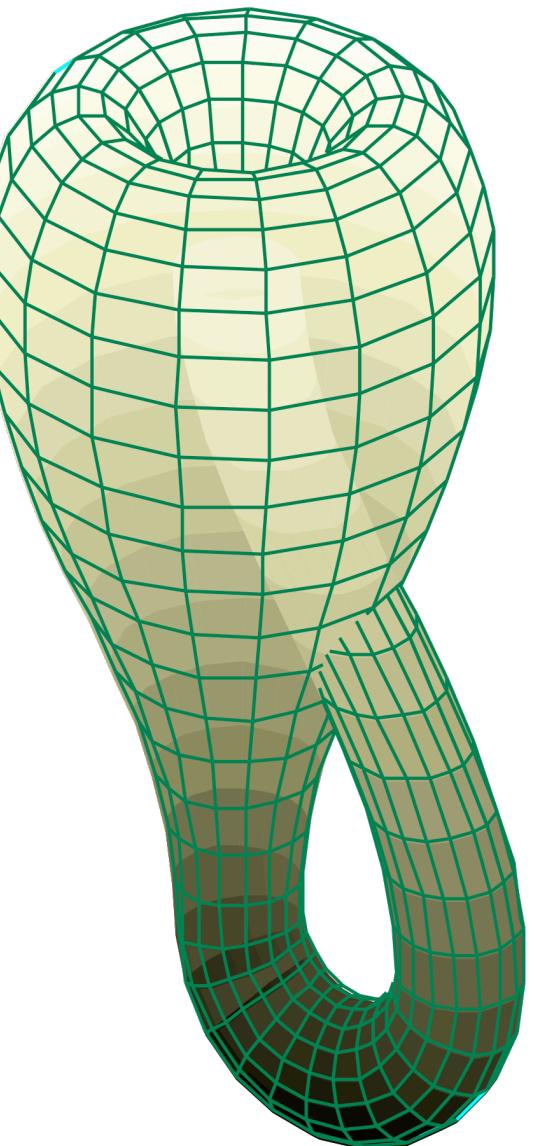


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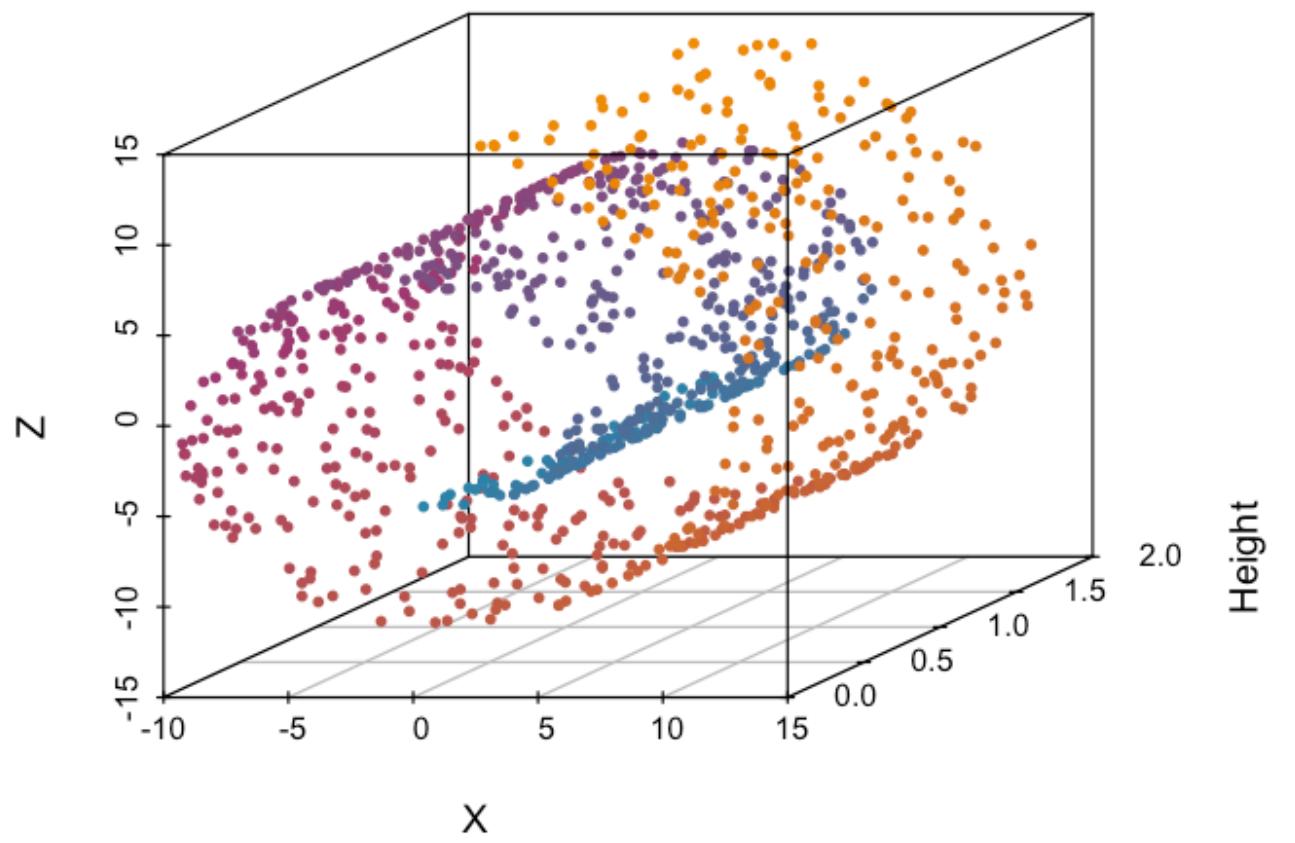


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  - Another example: a **rolled-up picture**

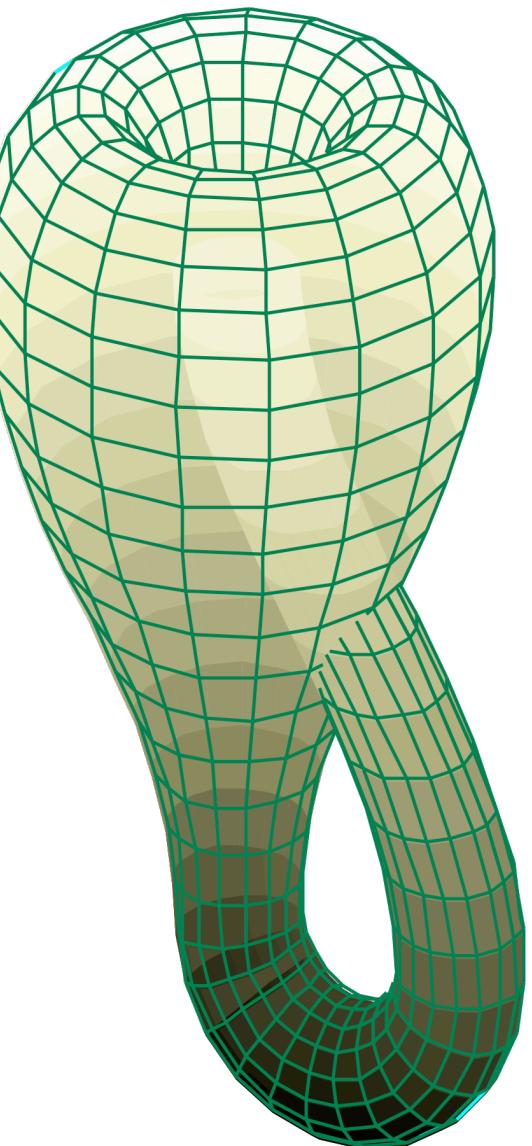


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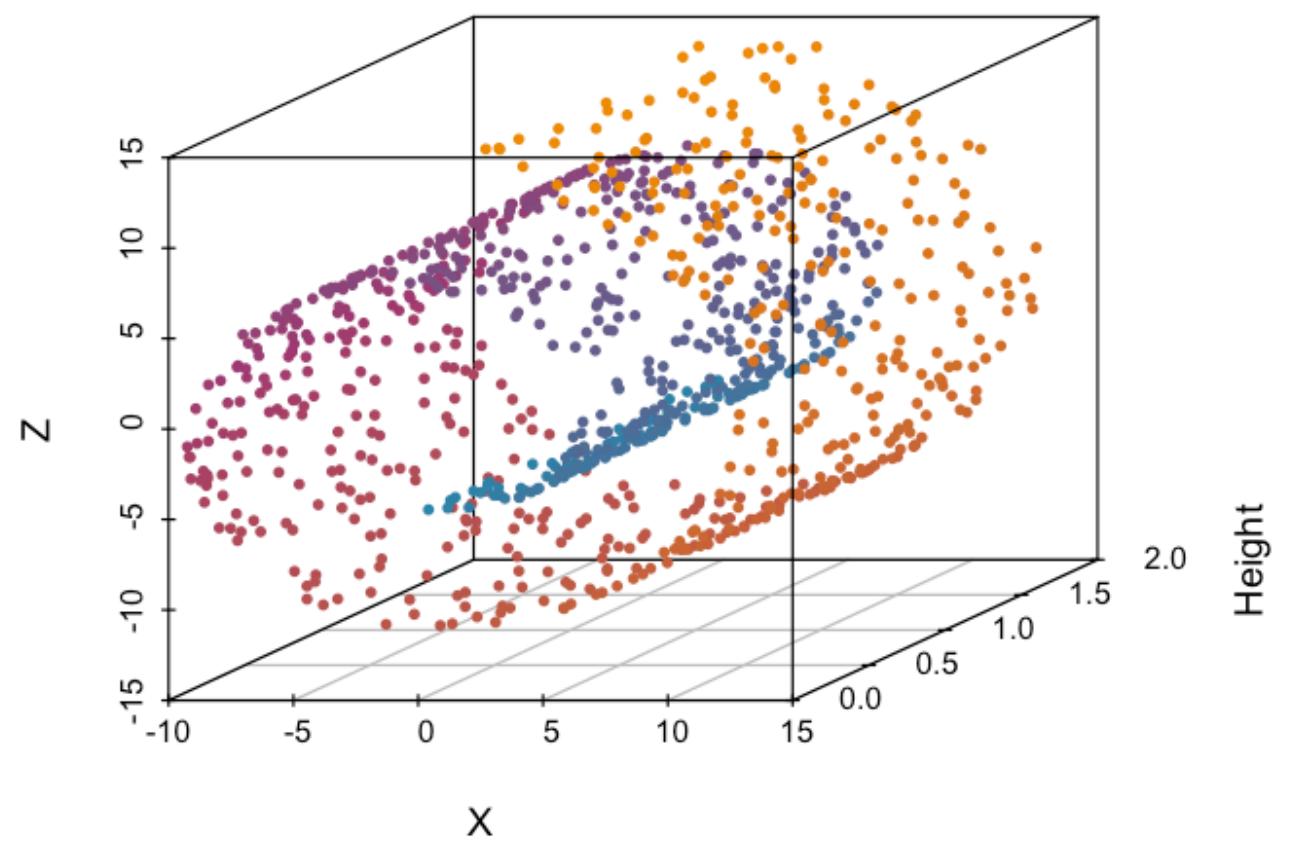


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  - Example: the **Earth's surface** can be **reasoned about in 2D** even though it curves
  - Another example: a **rolled-up picture**
- Goal: find the **dimensions that really matter**



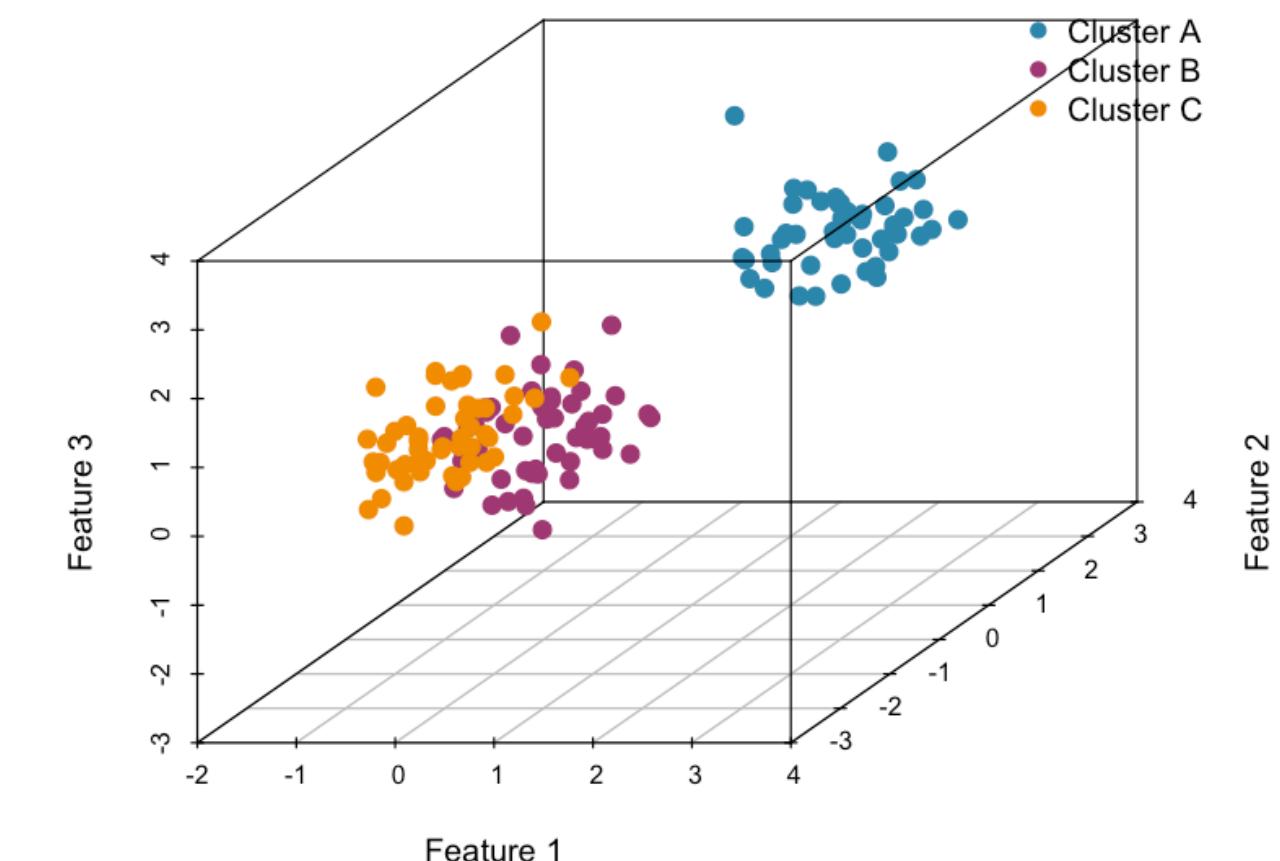
Rolled Up: A 2D Surface in 3D



# Principal Component Analysis (PCA)

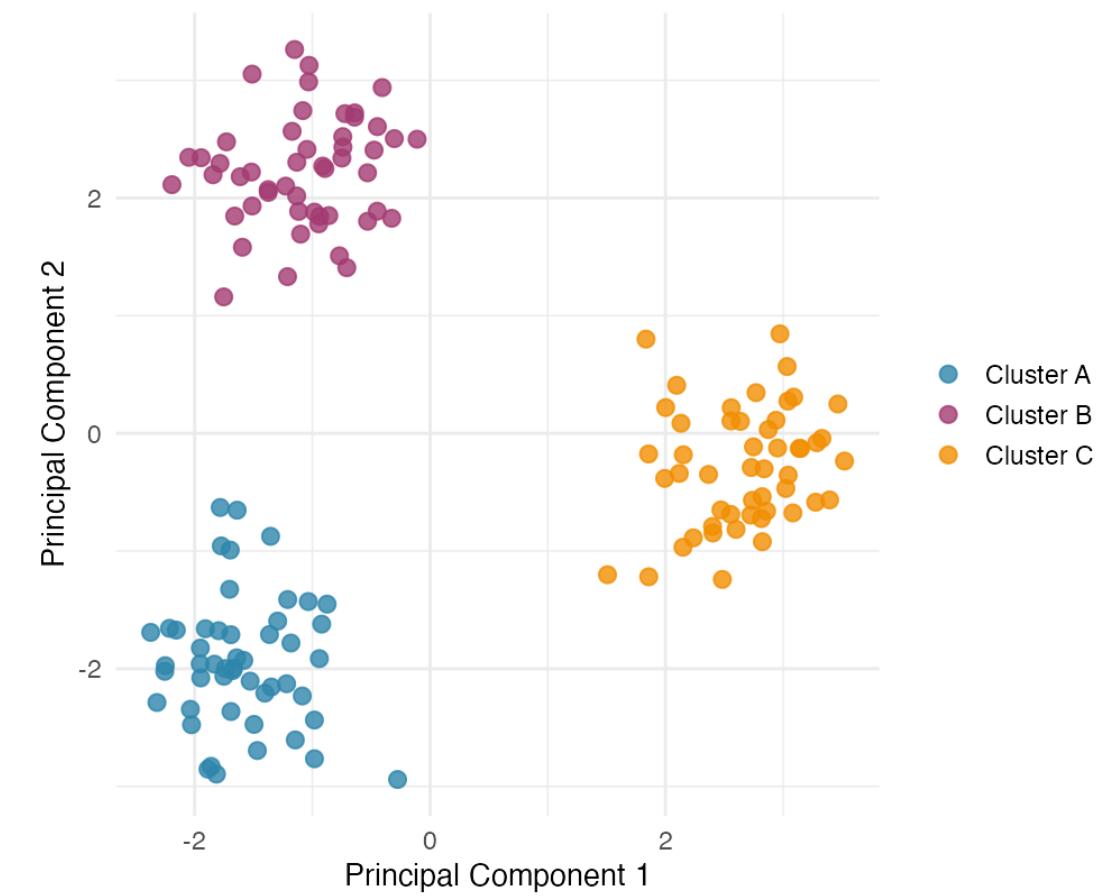
# PCA Generally

Three Clusters in 3D Space



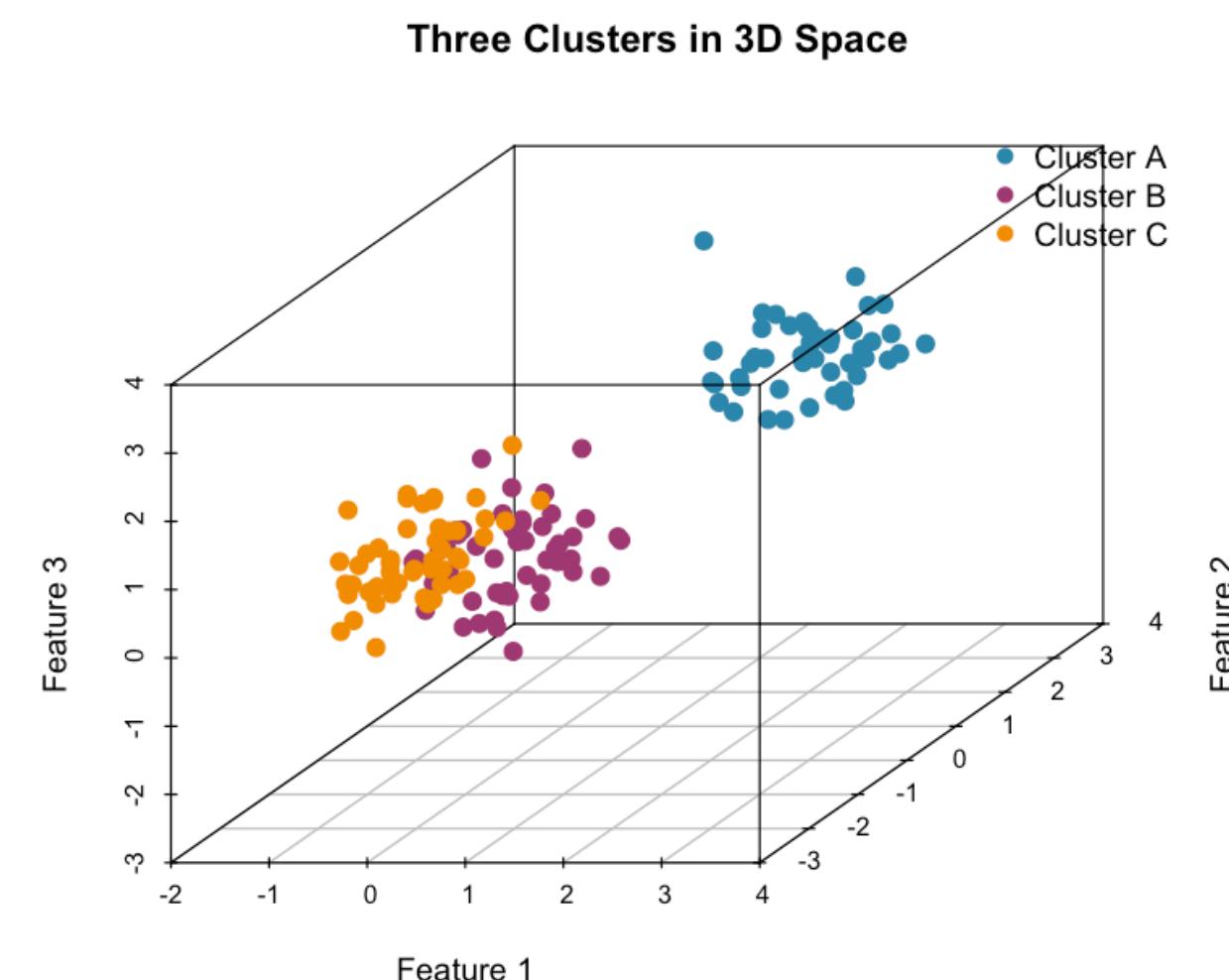
PCA Projection: 3D → 2D

PC1 explains 52.9% of variance, PC2 explains 44.1%

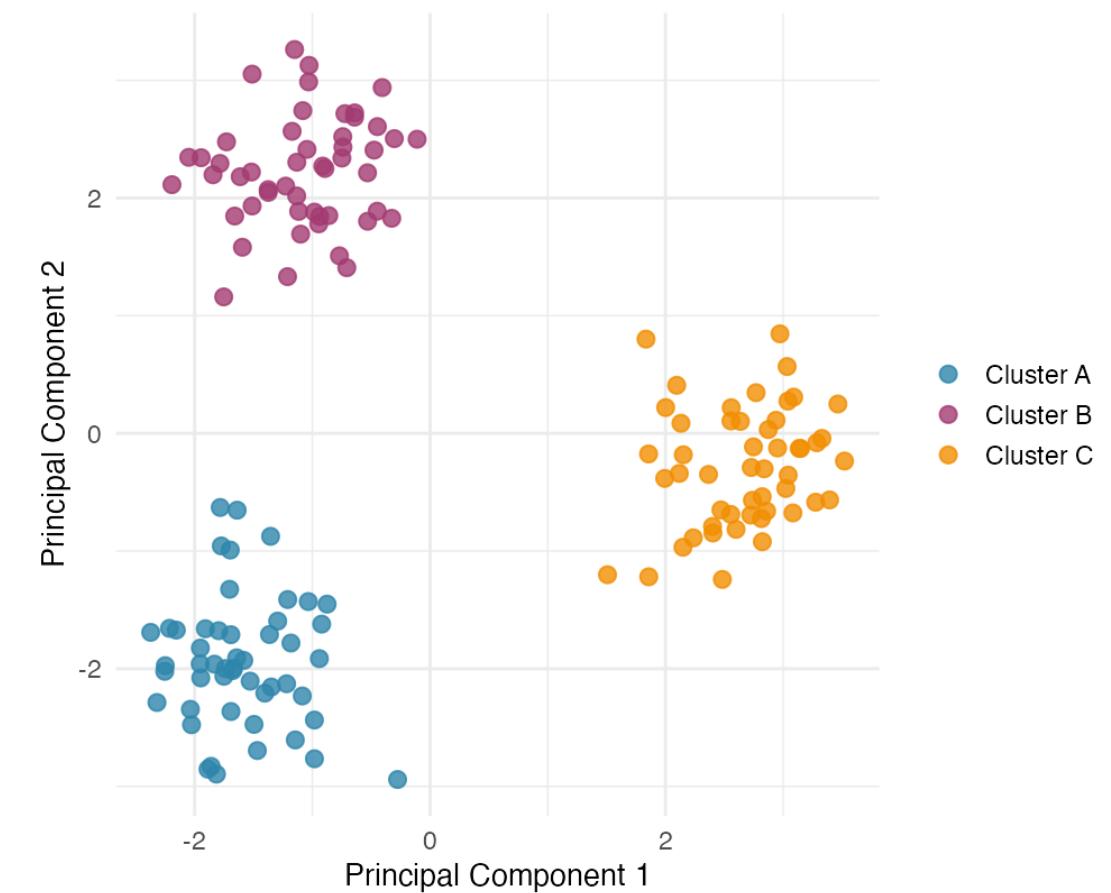


# PCA Generally

- Assume we have  $n$  datapoints of dimension  $d$ 
  - I.e. a matrix  $X$  of size  $d \times n$

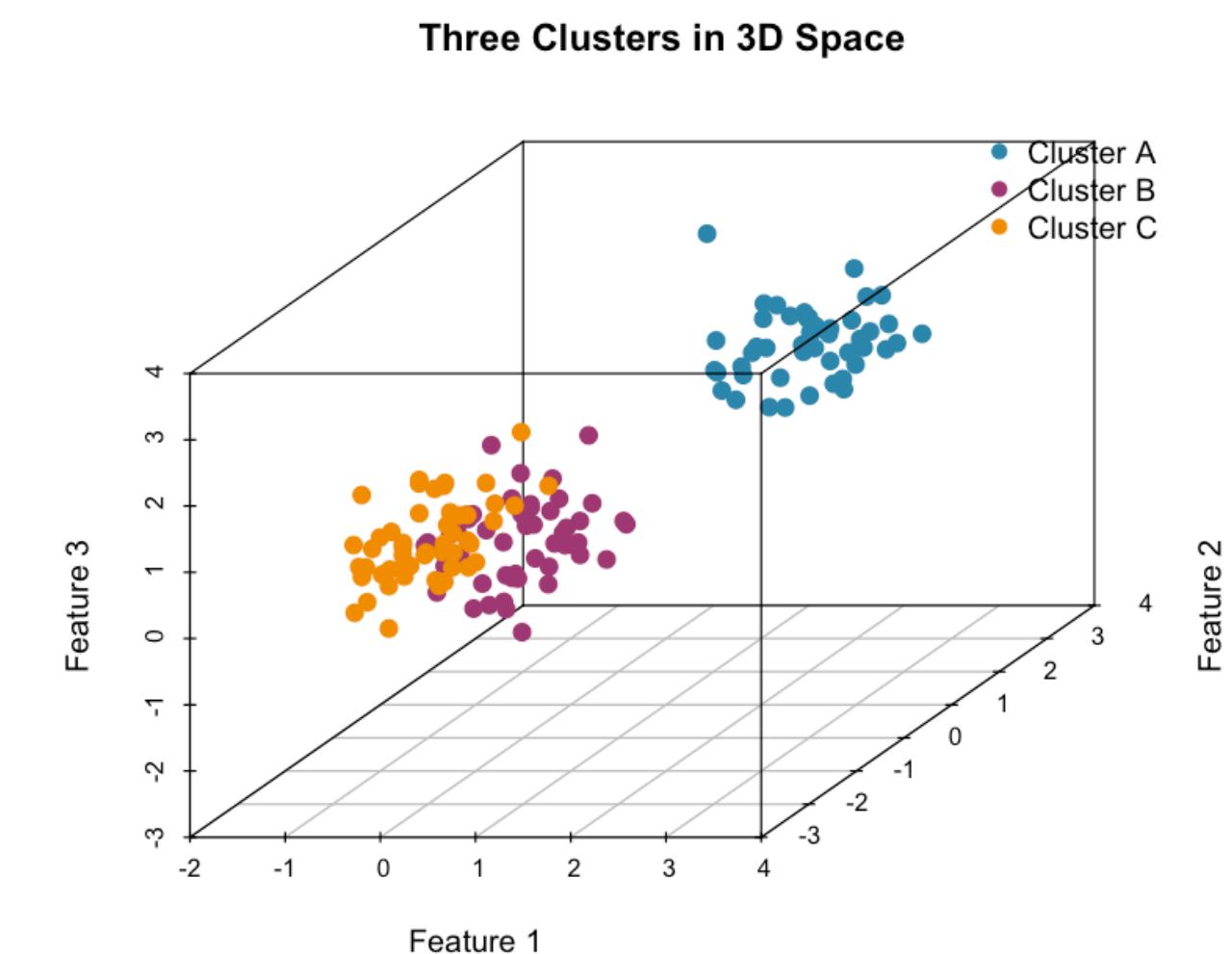


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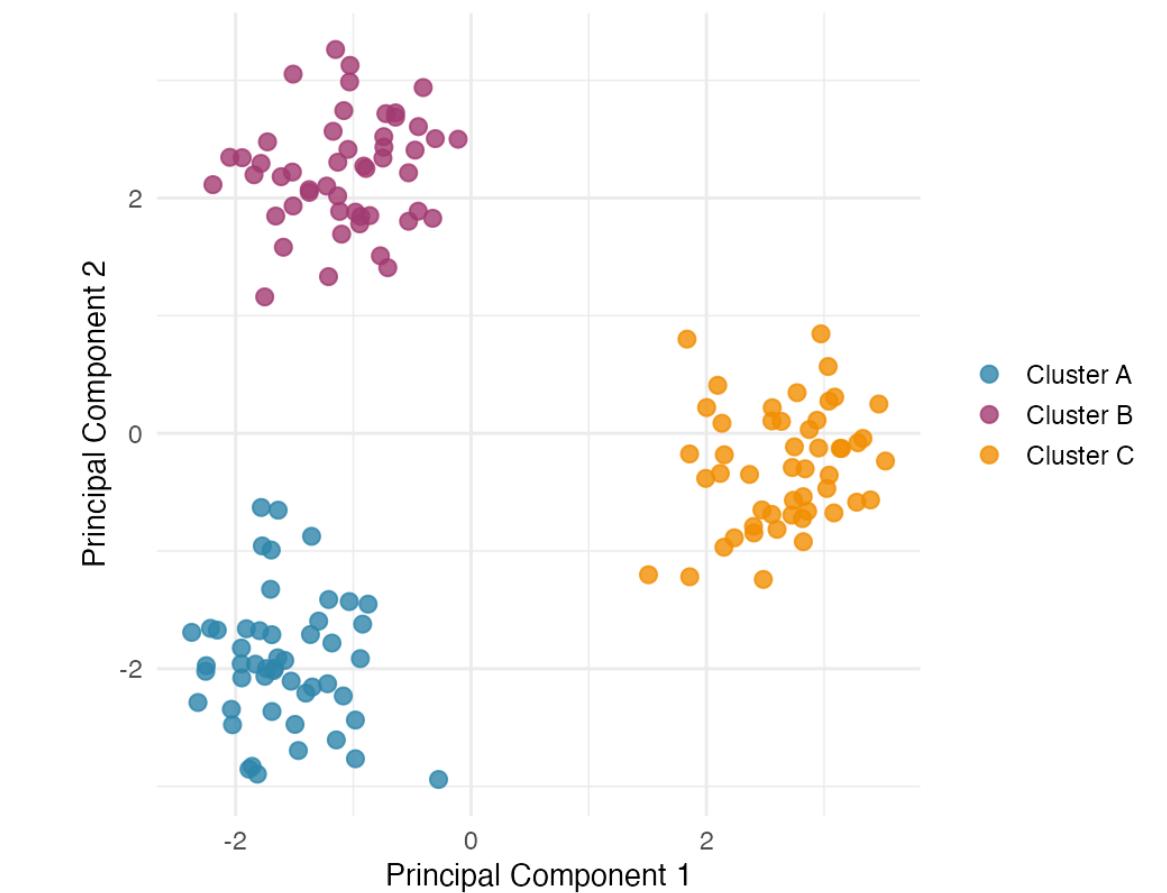


# PCA Generally

- Assume we have  $n$  datapoints of dimension  $d$ 
  - I.e. a matrix  $X$  of size  $d \times n$
- Goal: find the **most informative**  $k < d$  **dimensions** (the "Principal Components")
  - These are the directions that **explain the most variance** in the data
  - Put another way: the **axes** along which the **data varies the most**

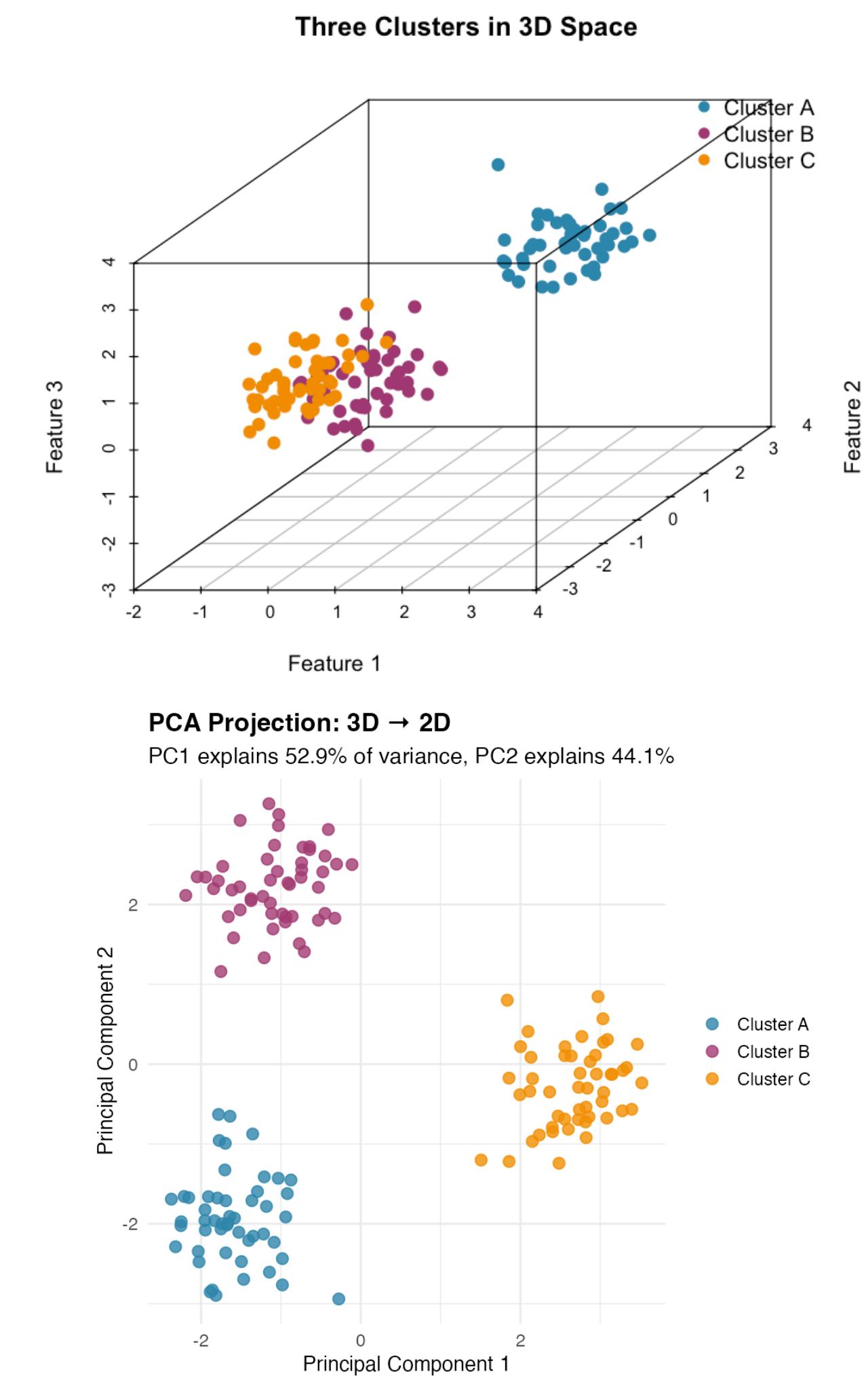


PCA Projection: 3D  $\rightarrow$  2D  
PC1 explains 52.9% of variance, PC2 explains 44.1%



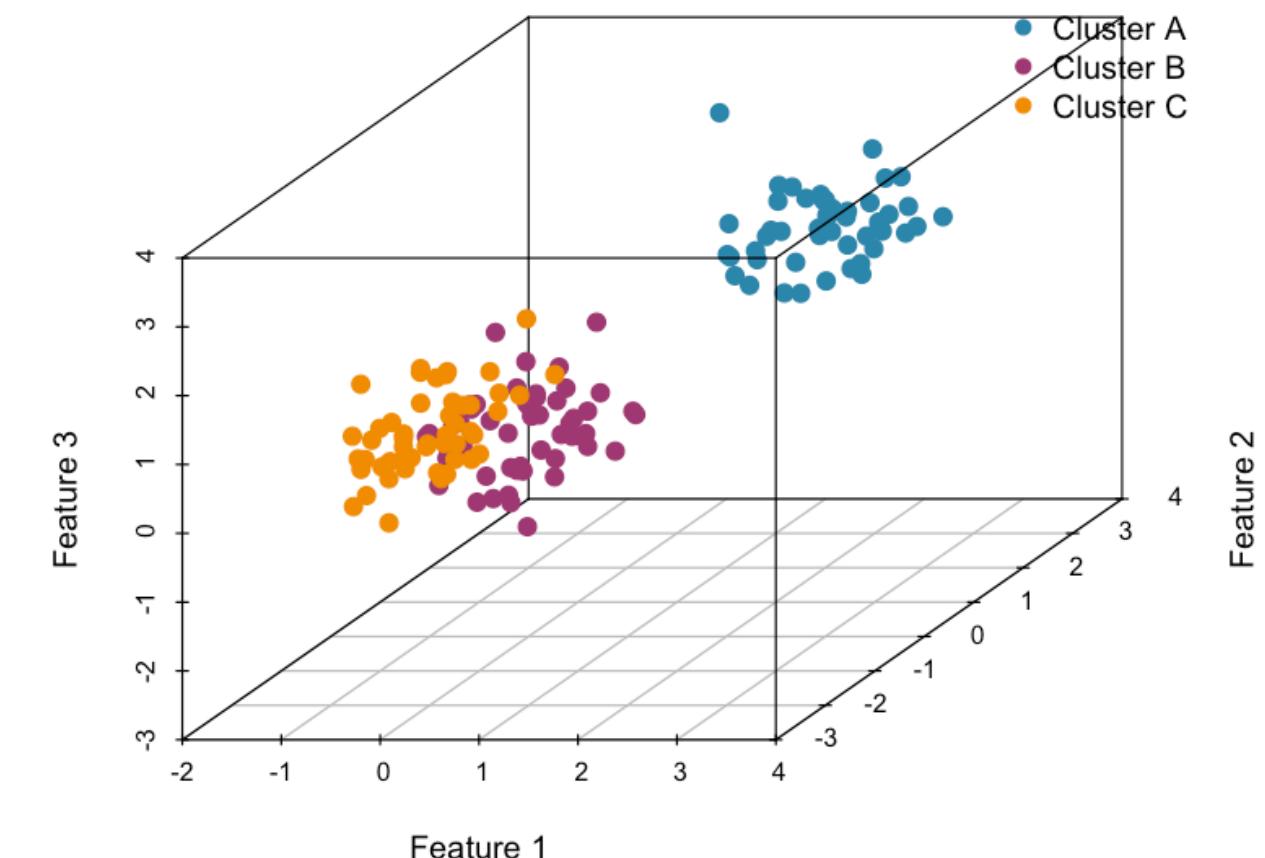
# PCA Generally

- Assume we have  $n$  datapoints of dimension  $d$ 
  - I.e. a matrix  $X$  of size  $d \times n$
- Goal: find the **most informative**  $k < d$  **dimensions** (the "Principal Components")
  - These are the directions that **explain the most variance** in the data
  - Put another way: the **axes** along which the **data varies the most**
- Then **project** onto the **(hyper-)plane formed by those axes**
  - Projection  $\approx$  **casting a shadow onto**



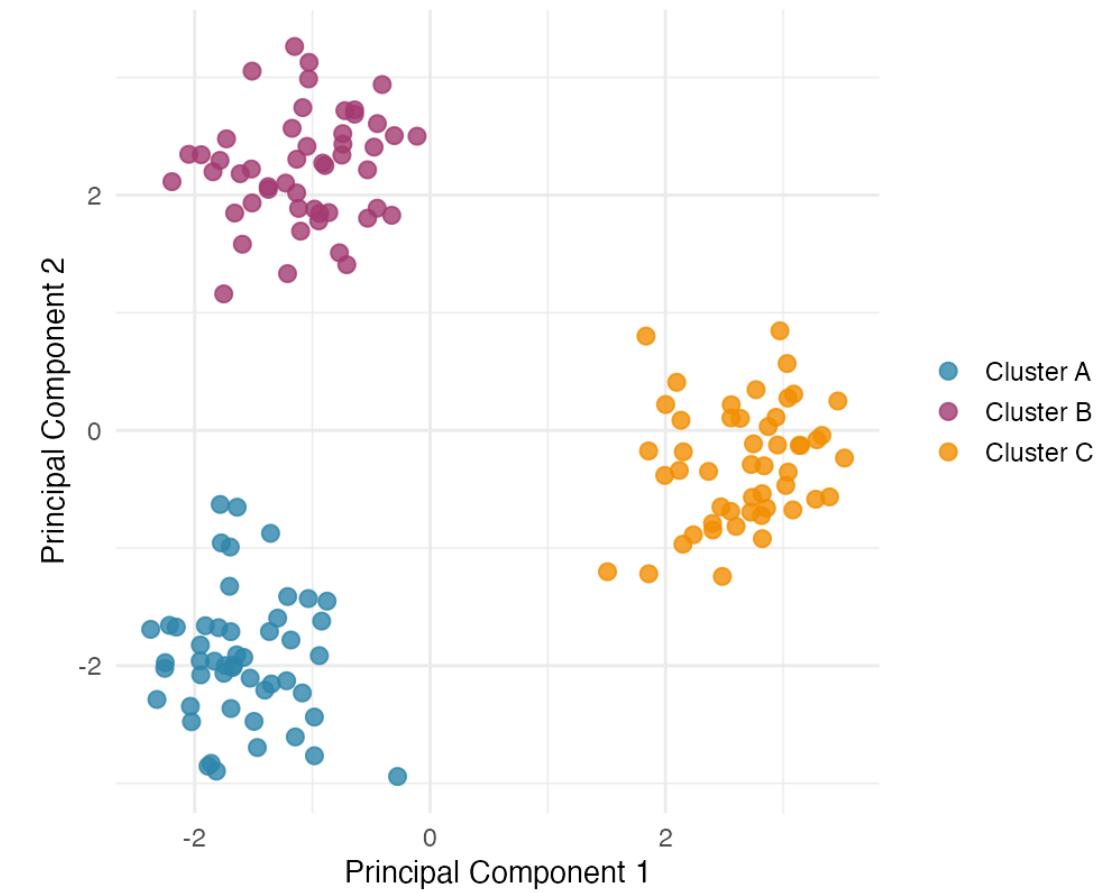
# PCA and Manifolds

Three Clusters in 3D Space



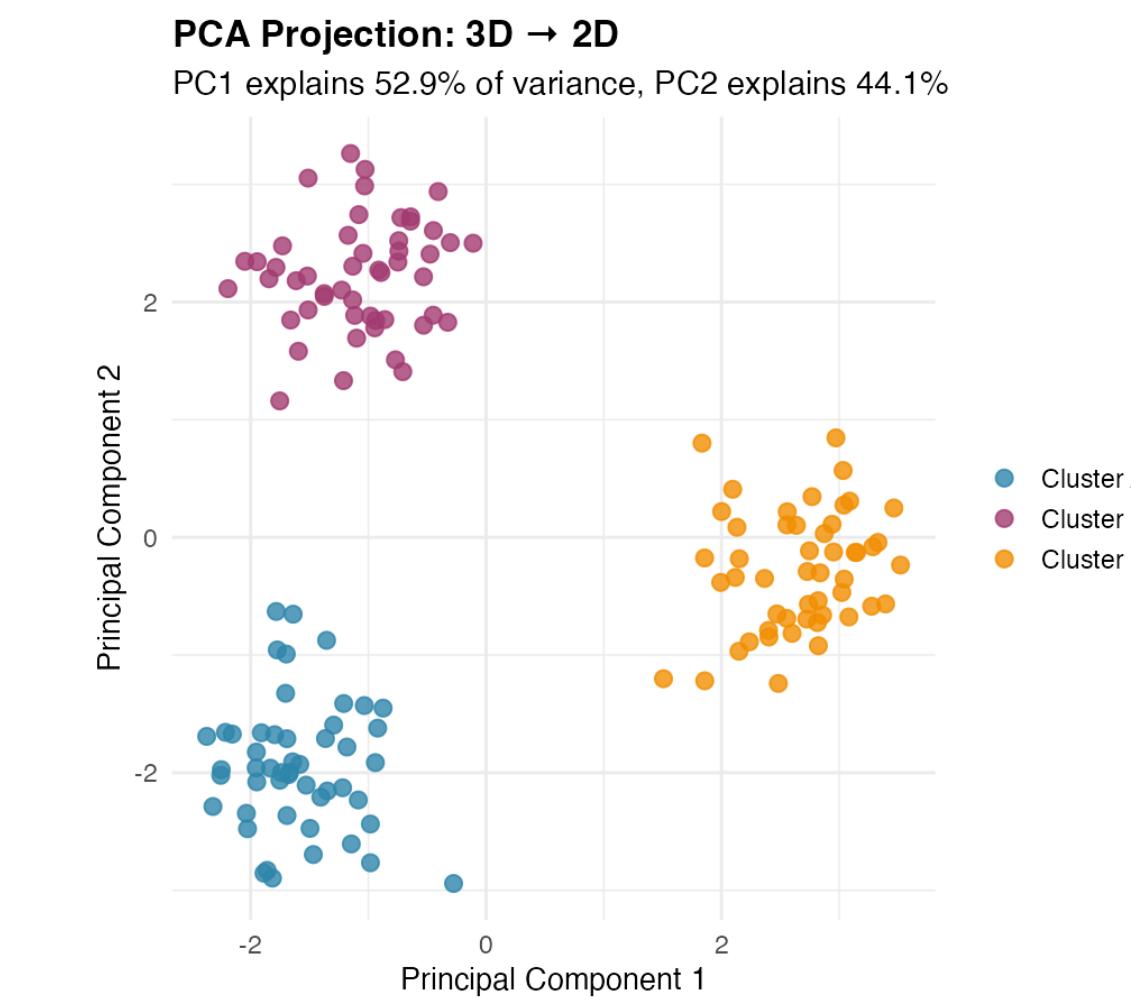
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# PCA and Manifolds

- Assumes the data is **near a linear manifold** (aka a [hyper-]plane)
  - Treats other dimensions as **noise**
  - Plane treated as the "**true nature**" of the data

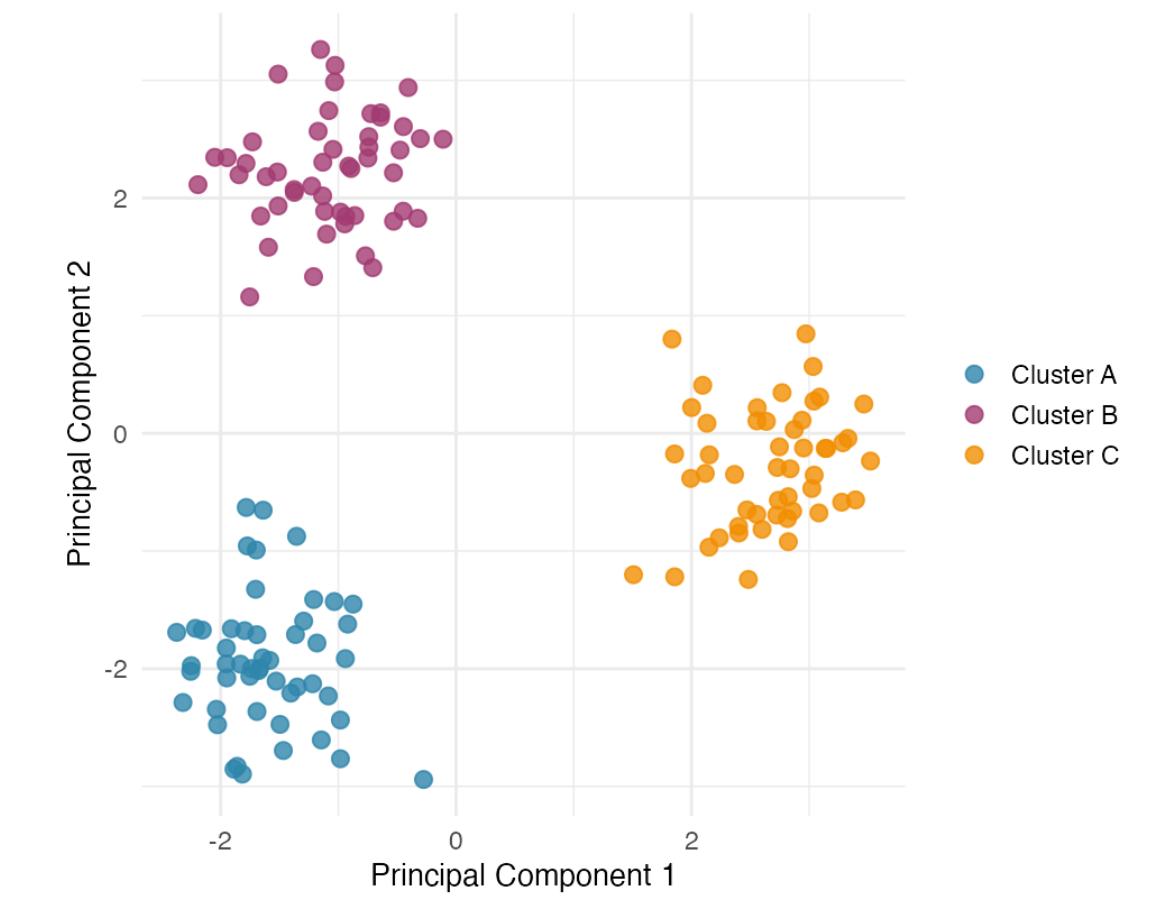


# PCA and Manifolds

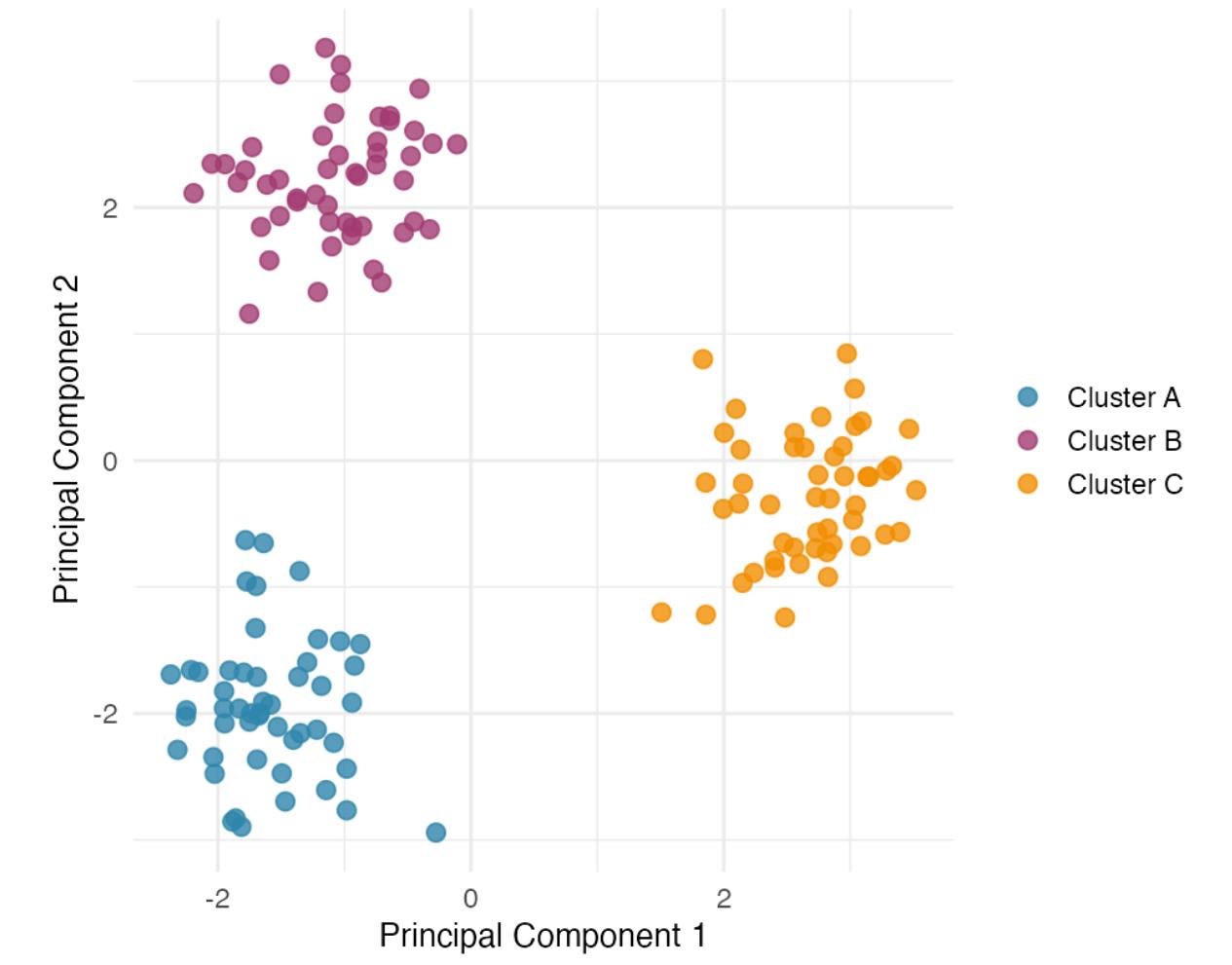
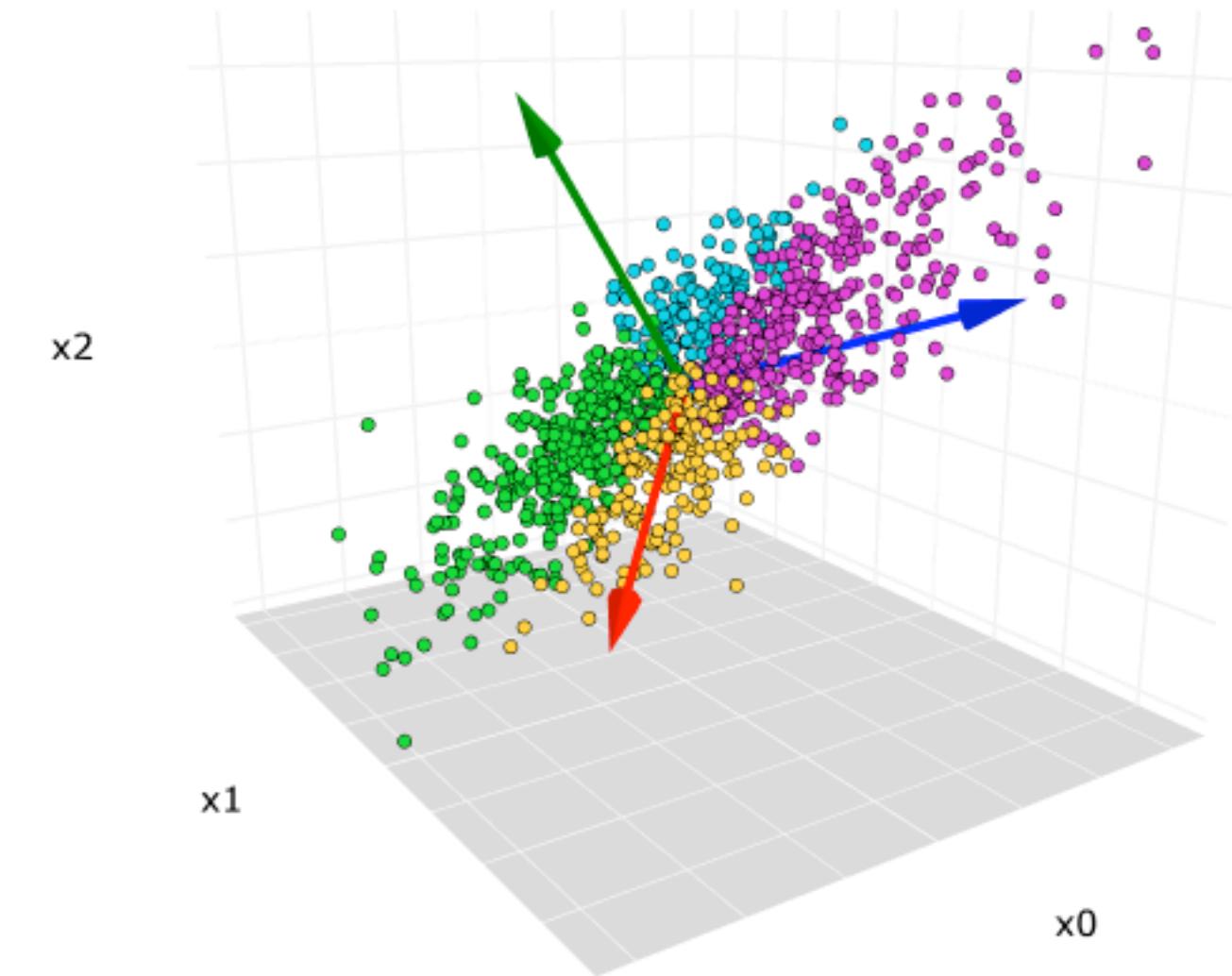
- Assumes the data is **near a linear manifold** (aka a [hyper-]plane)
  - Treats other dimensions as **noise**
  - Plane treated as the "**true nature**" of the data
- This is an **inductive bias!**
  - We'll see how this can **fail** (non-linear manifolds)



PCA Projection: 3D → 2D  
PC1 explains 52.9% of variance, PC2 explains 44.1%

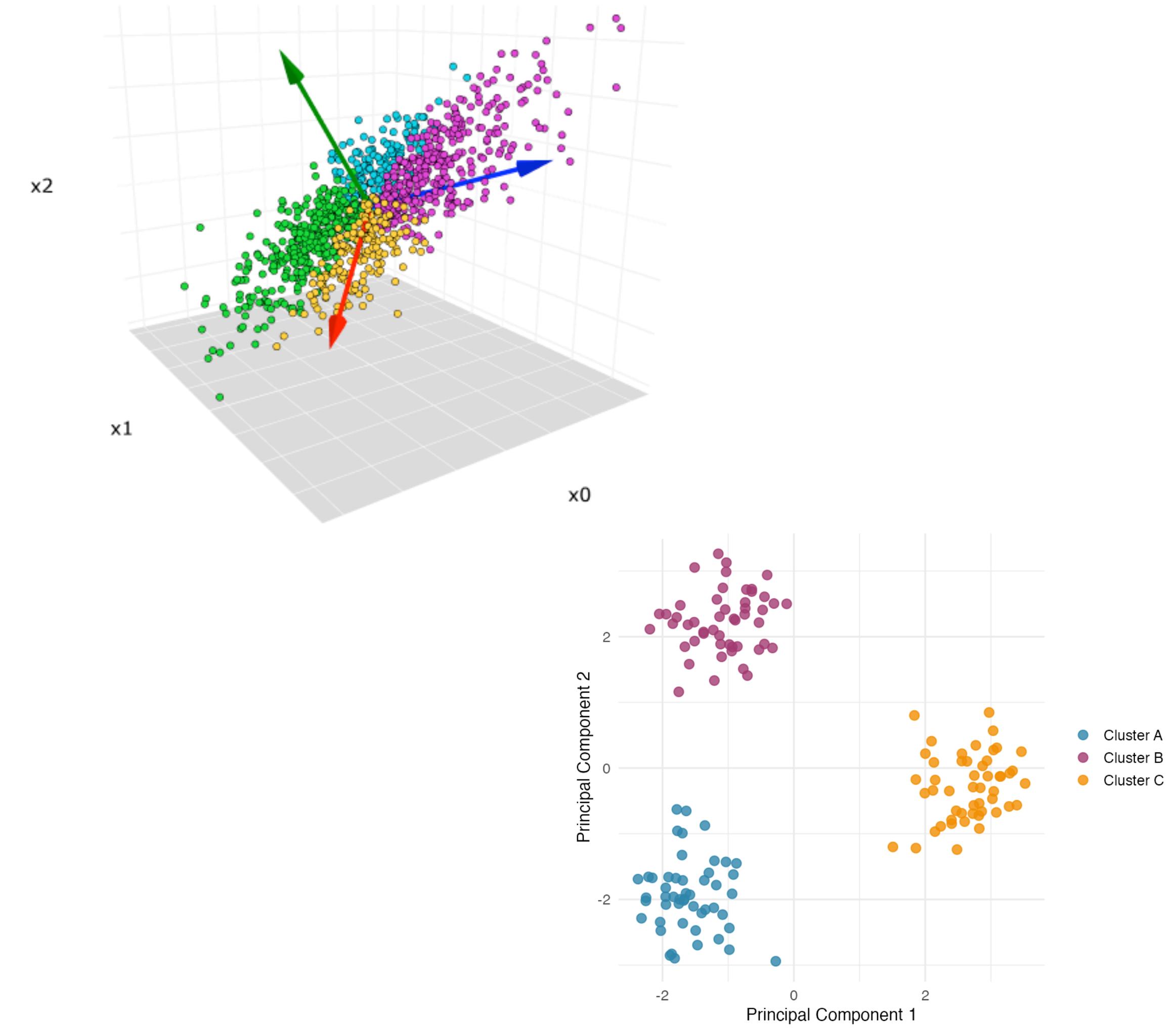


# PCA Algorithm



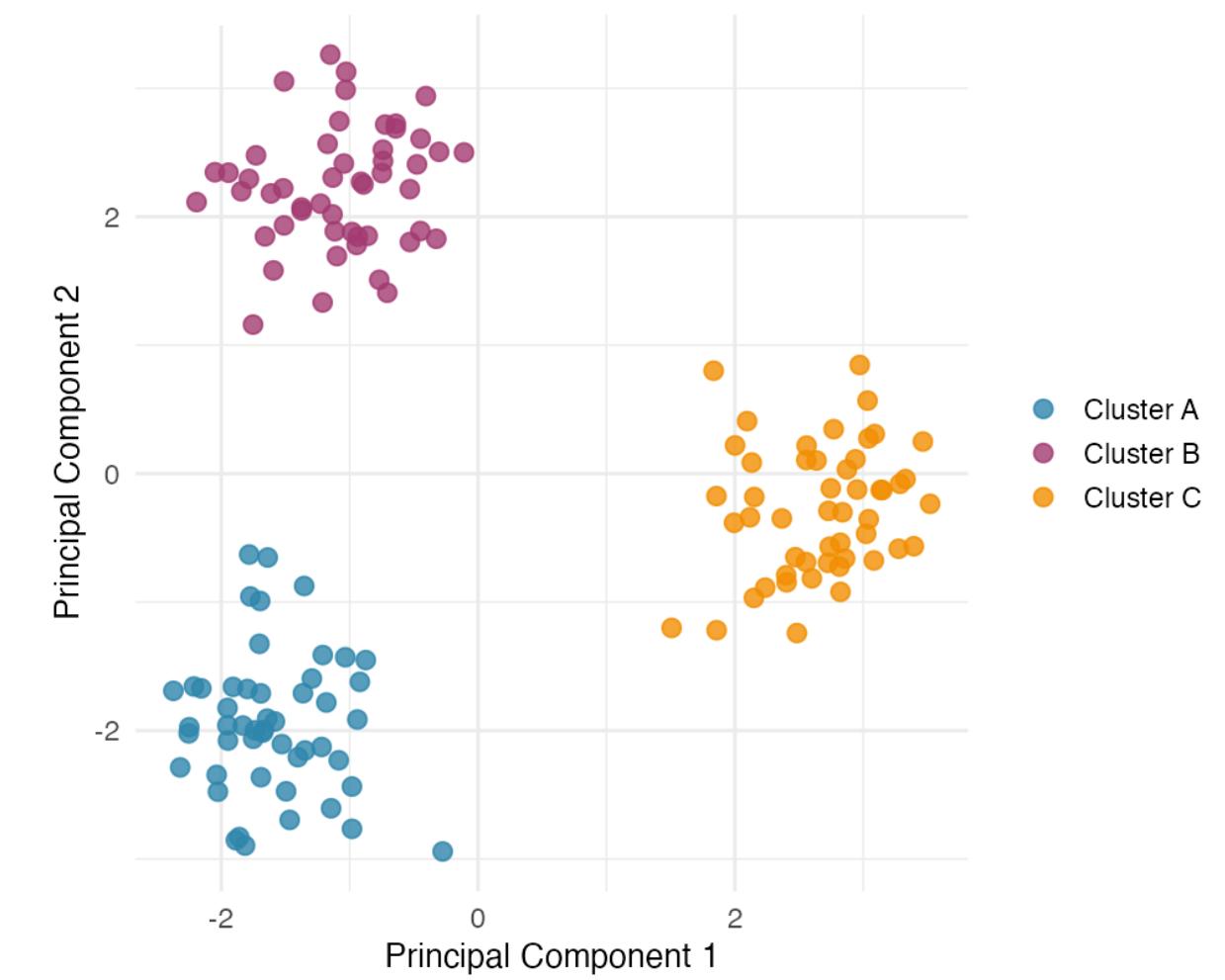
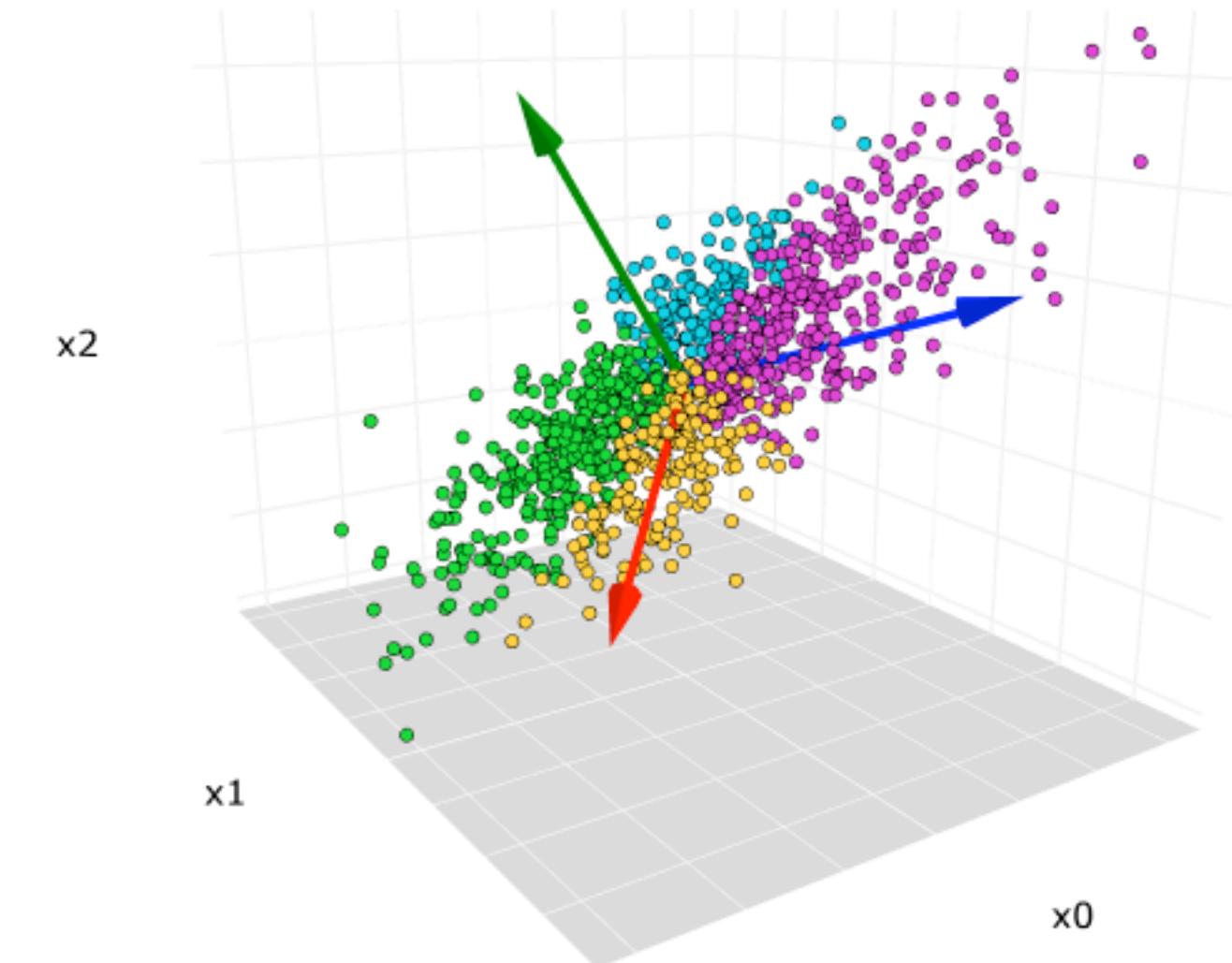
# PCA Algorithm

- Find the  $k$  **directions of greatest variance**
  - Gotten from the **covariance matrix** of the data (next slide)
  - These will be **orthogonal**



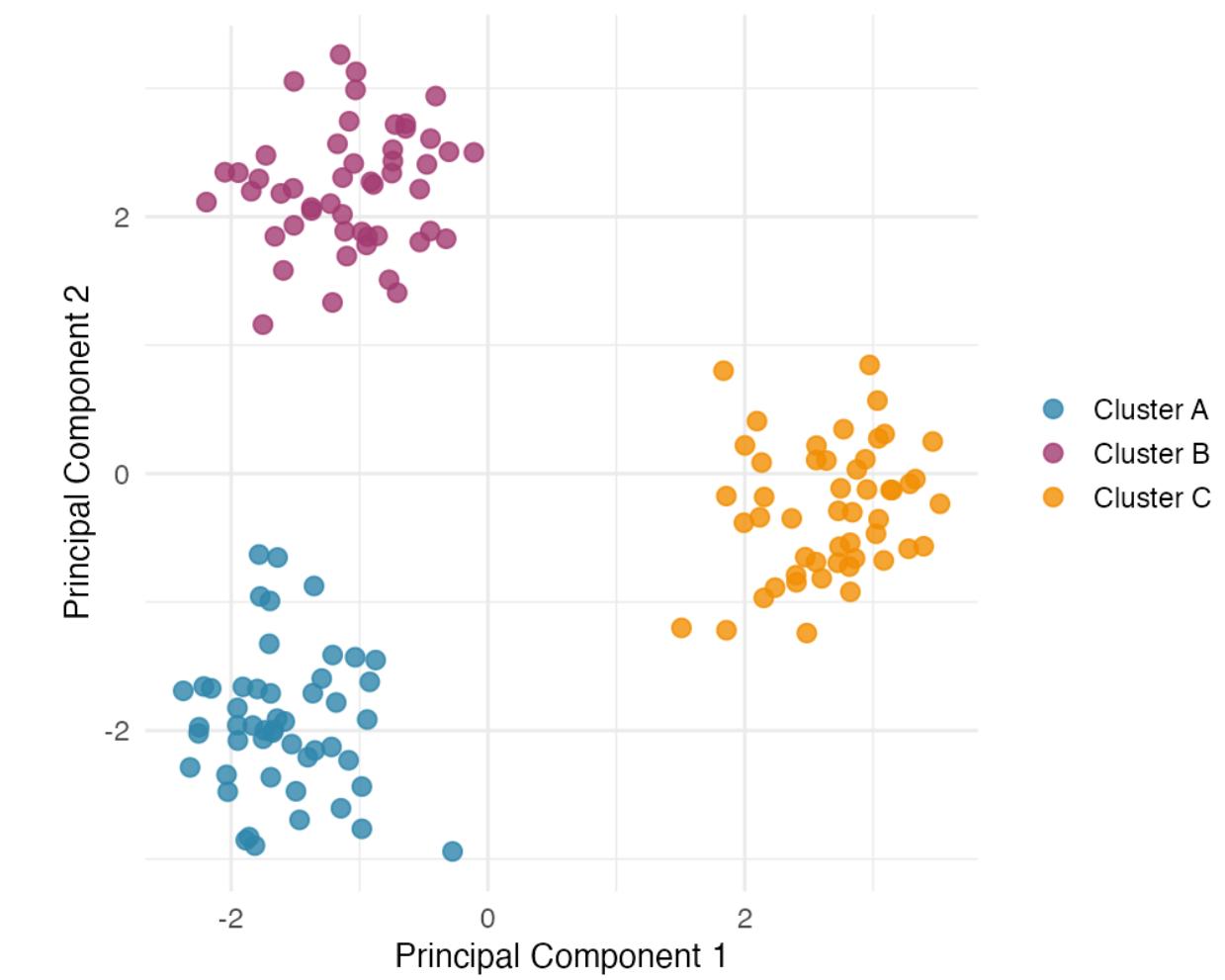
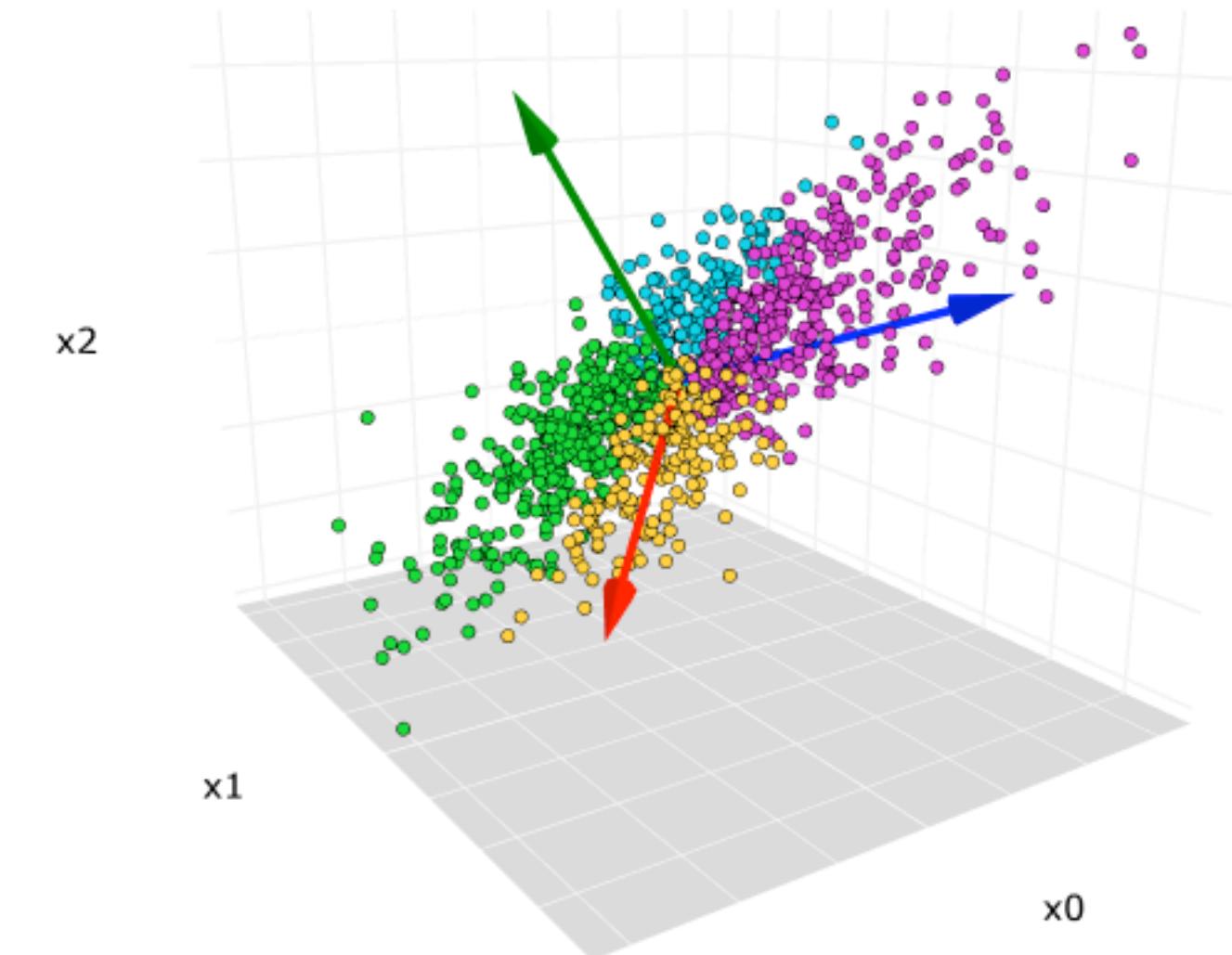
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# PCA Algorithm

- Find the  $k$  **directions of greatest variance**
  - Gotten from the **covariance matrix** of the data (next slide)
  - These will be **orthogonal**
- The direction vectors **define a plane**
- **Project the data onto that plane**
  - This is fairly simple Linear Algebra



# Covariance Matrix

$$\sum_k \mathbf{x}_k \mathbf{x}_k^T = \sum_k \begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ x_d \end{bmatrix} [x_1 \ x_2 \ \cdots \ x_d] = \sum_k \begin{bmatrix} x_1 x_1 & x_1 x_2 & \cdots & x_1 x_d \\ x_2 x_1 & x_2 x_2 & \cdots & x_2 x_d \\ \vdots & \vdots & \ddots & \vdots \\ x_d x_1 & x_d x_2 & \cdots & x_d x_d \end{bmatrix}$$

$$\mathbf{X}^T \mathbf{X} = \begin{bmatrix} | & | & & | \\ \mathbf{x}_1 & \mathbf{x}_2 & \cdots & \mathbf{x}_n \\ | & | & & | \end{bmatrix} \begin{bmatrix} \cdots & \mathbf{x}_1^T & \cdots \\ \cdots & \mathbf{x}_2^T & \cdots \\ \vdots & & \vdots \\ \cdots & \mathbf{x}_n^T & \cdots \end{bmatrix}$$

$$\Sigma = \begin{bmatrix} \sigma_1^2 & \sigma_{12} & \cdots & \sigma_{1d} \\ \sigma_{21} & \sigma_2^2 & \cdots & \sigma_{2d} \\ \vdots & \vdots & \ddots & \vdots \\ \sigma_{d1} & \sigma_{d2} & \cdots & \sigma_d^2 \end{bmatrix}$$

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one high-dimensional datapoint

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# Covariance Matrix

These three definitions are equivalent!

$$\sum_k \mathbf{x}_k \mathbf{x}_k^T = \sum_k \begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ x_d \end{bmatrix} [x_1 \ x_2 \ \cdots \ x_d] = \sum_k \begin{bmatrix} x_1 x_1 & x_1 x_2 & \cdots & x_1 x_d \\ x_2 x_1 & x_2 x_2 & \cdots & x_2 x_d \\ \vdots & \vdots & \ddots & \vdots \\ x_d x_1 & x_d x_2 & \cdots & x_d x_d \end{bmatrix}$$

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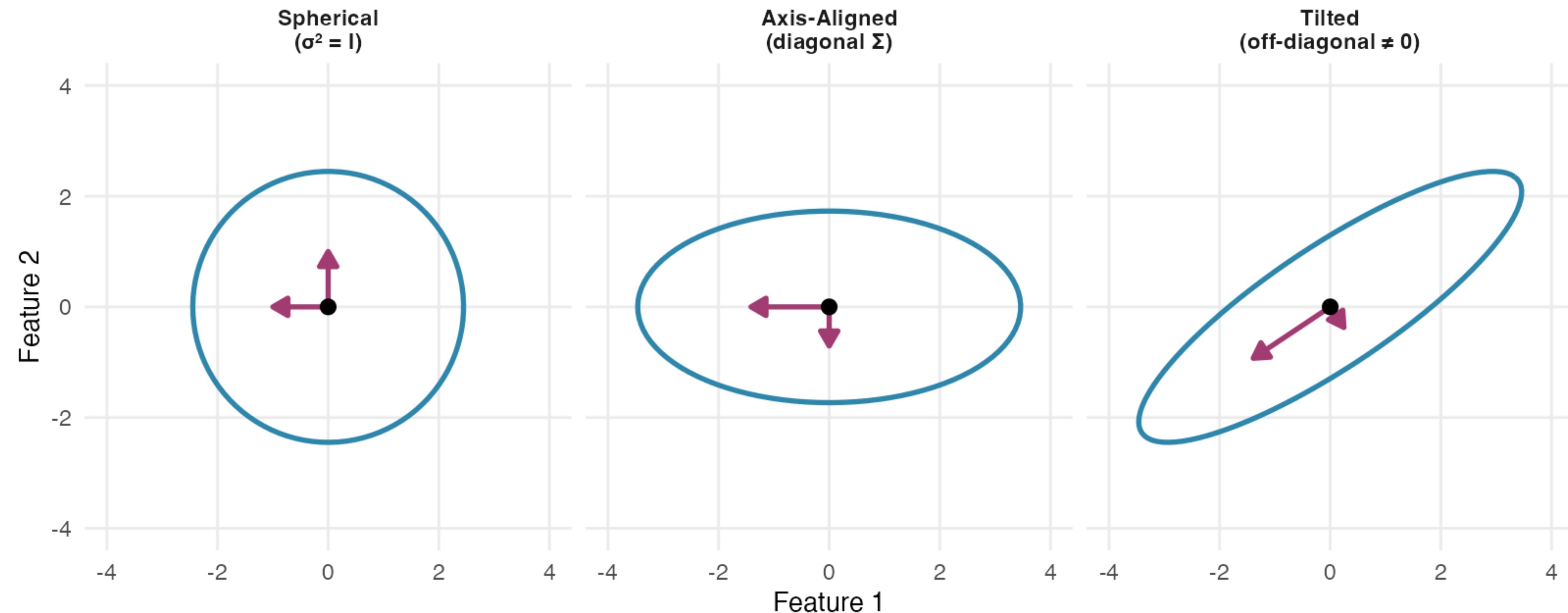
How much **co-variation** does each pair of variables have?

$$\Sigma = \begin{bmatrix} \sigma_1^2 & \sigma_{12} & \cdots & \sigma_{1d} \\ \sigma_{21} & \sigma_2^2 & \cdots & \sigma_{2d} \\ \vdots & \vdots & \ddots & \vdots \\ \sigma_{d1} & \sigma_{d2} & \cdots & \sigma_d^2 \end{bmatrix}$$

# Covariance matrix defines an ellipse

## Covariance Matrices as Ellipses

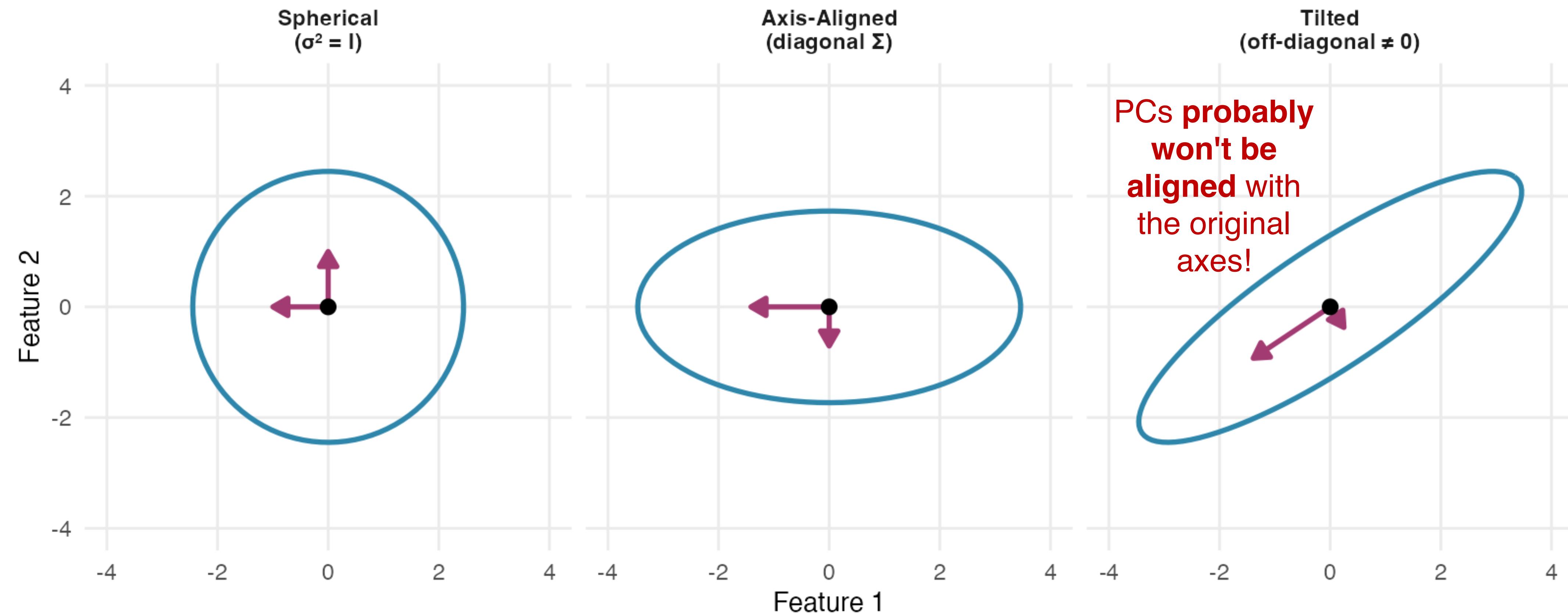
Arrows show eigenvectors  $\times \text{sqrt}(\text{eigenvalues})$  — the principal axes



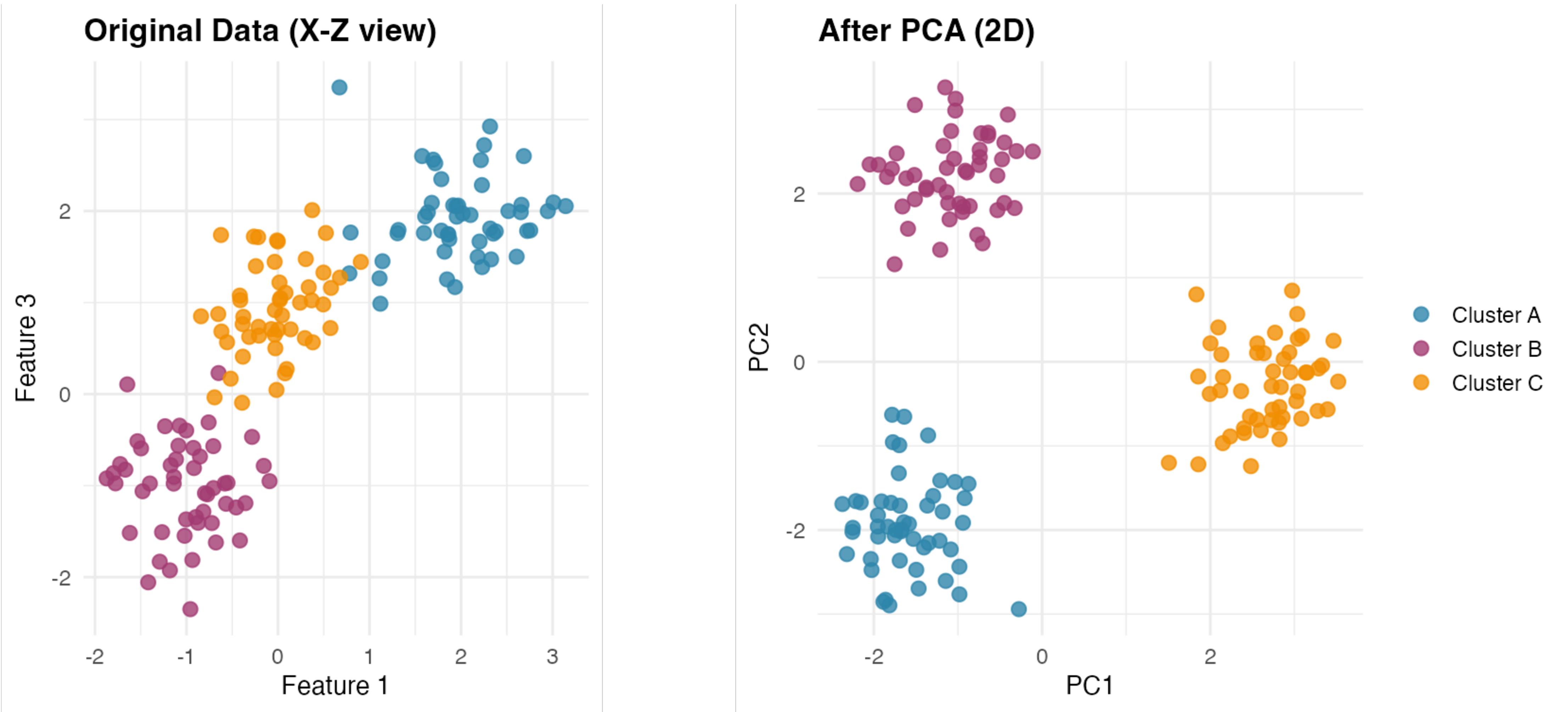
# Covariance matrix defines an ellipse

## Covariance Matrices as Ellipses

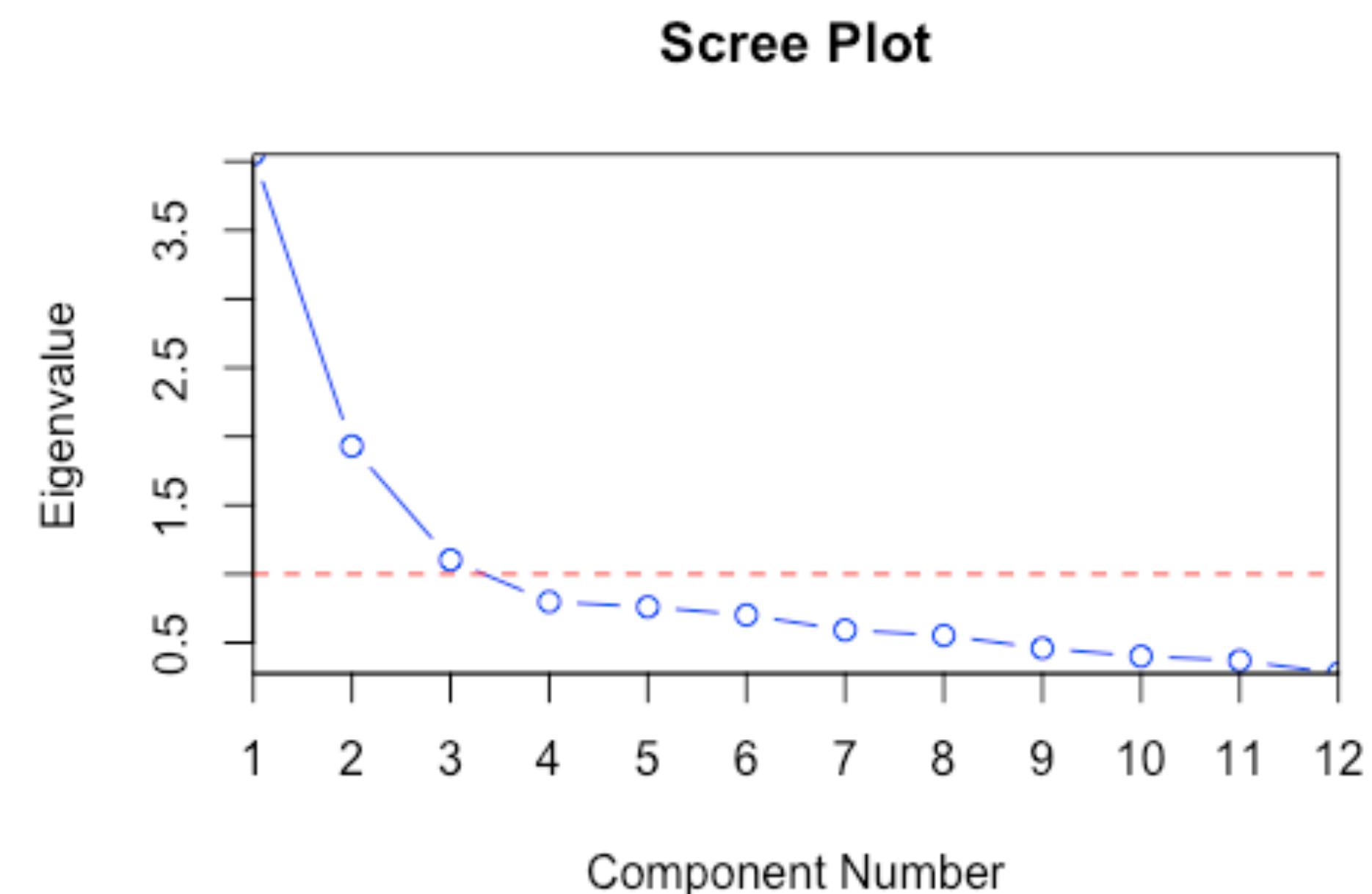
Arrows show eigenvectors  $\times \text{sqrt}(\text{eigenvalues})$  — the principal axes



# PCA aligns to PC axes

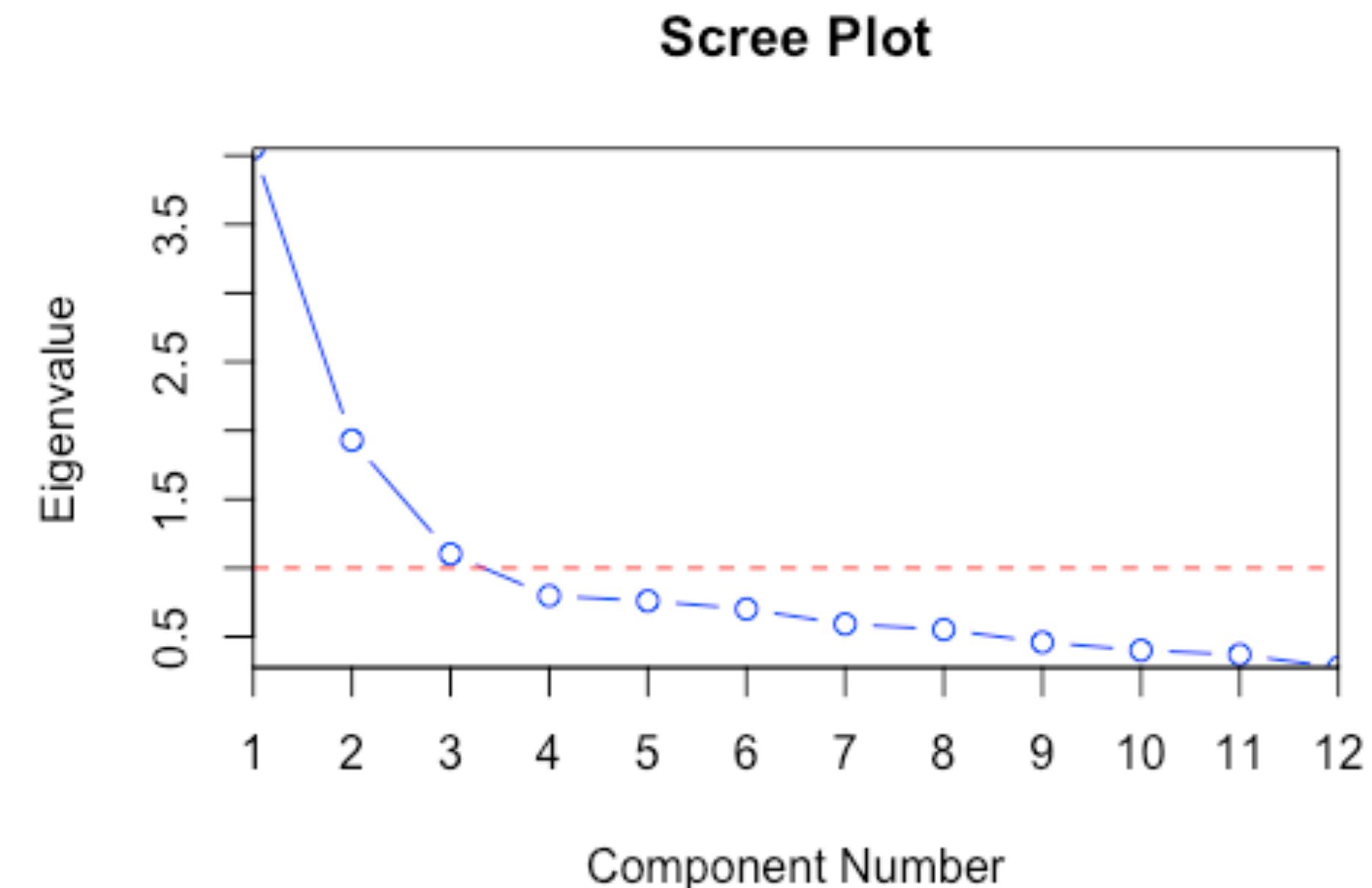


# Choosing Component Numbers



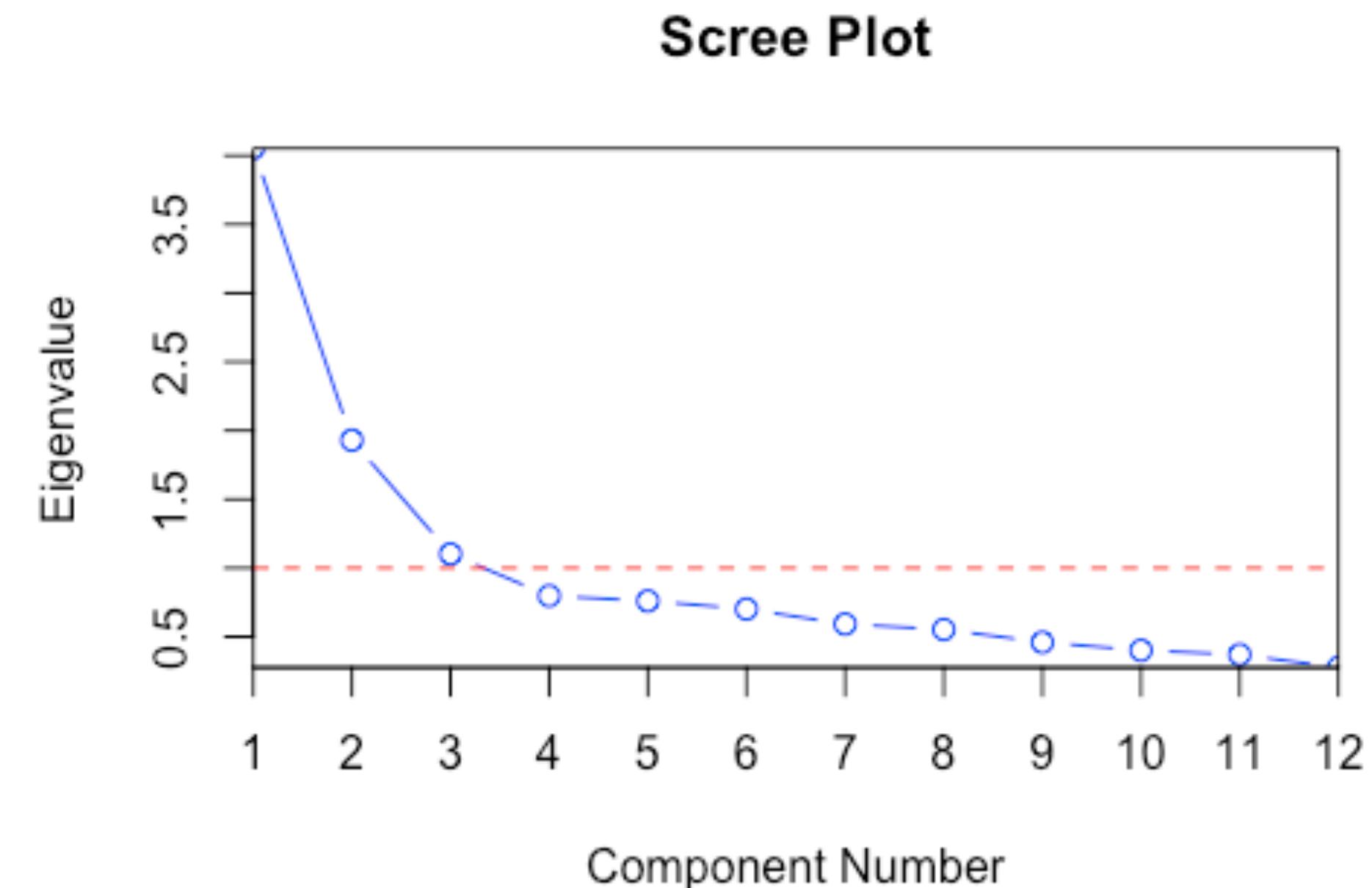
# Choosing Component Numbers

- Number of components  $k$  to keep usually chosen with a **Scree Plot**
  - Plot of **how much variance** each component explains
  - These are also called the **Eigenvalues**

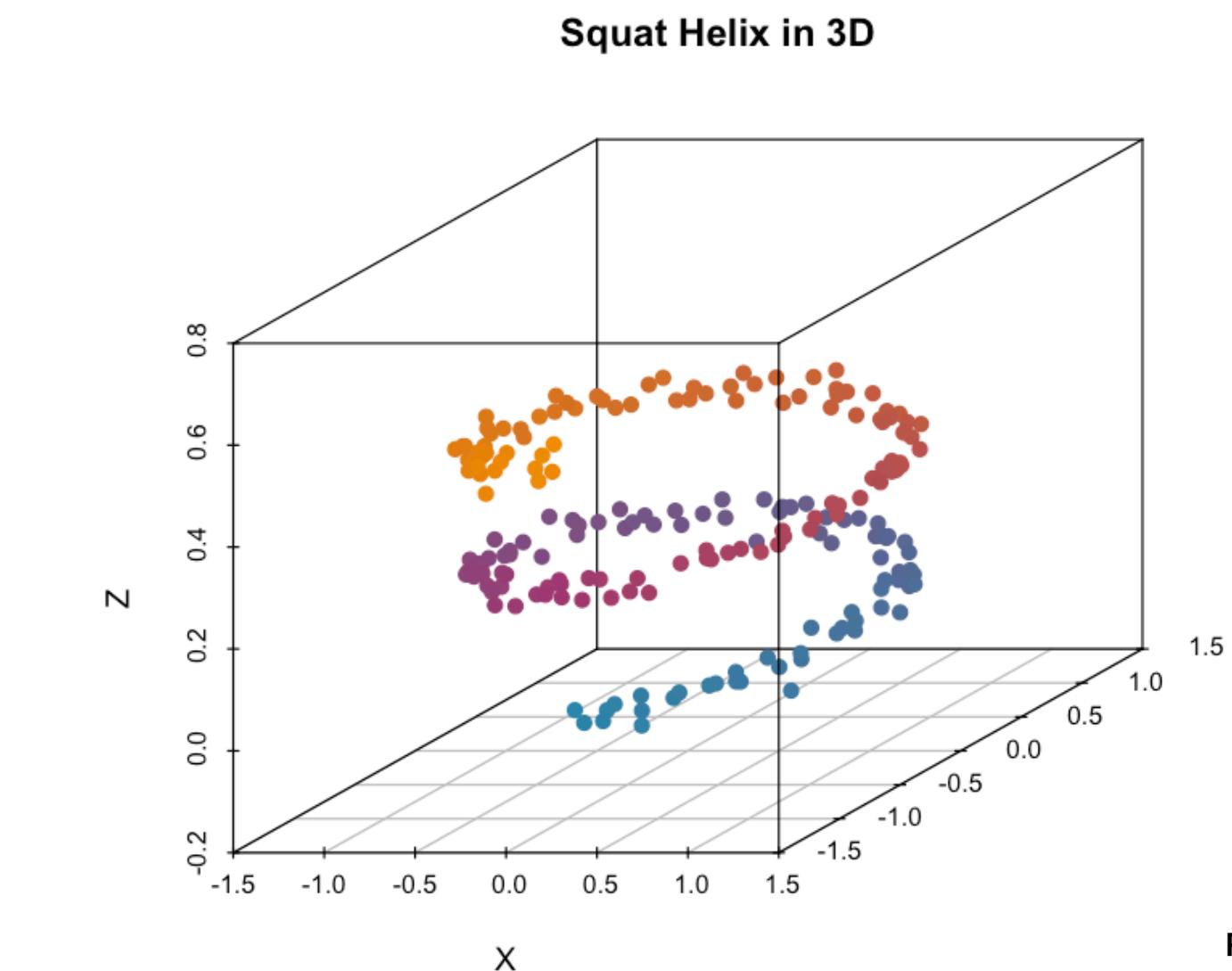


# Choosing Component Numbers

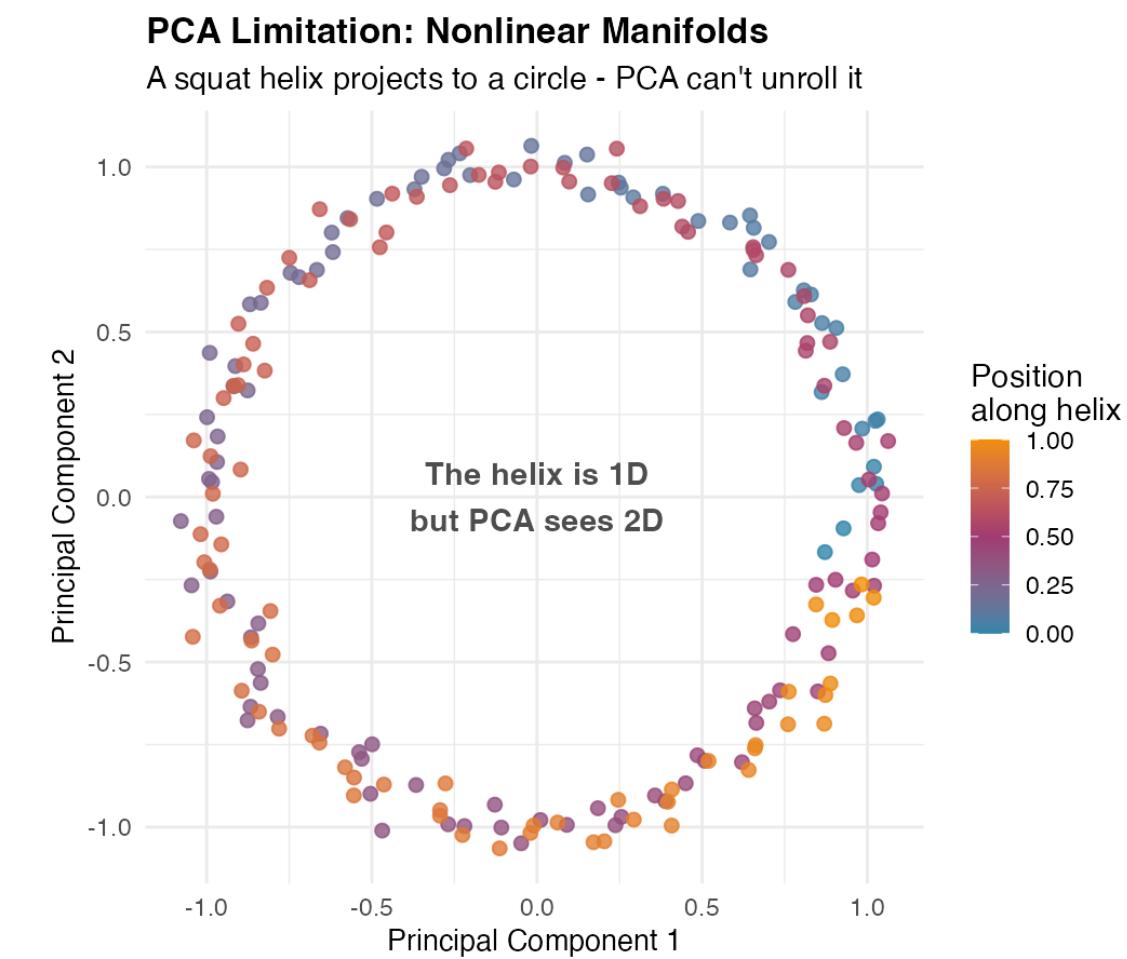
- Number of components  $k$  to keep usually chosen with a **Scree Plot**
  - Plot of **how much variance** each component explains
  - These are also called the **Eigenvalues**
- Often the **point of inflection** is chosen
  - (i.e. diminishing returns)



# Where PCA Fails

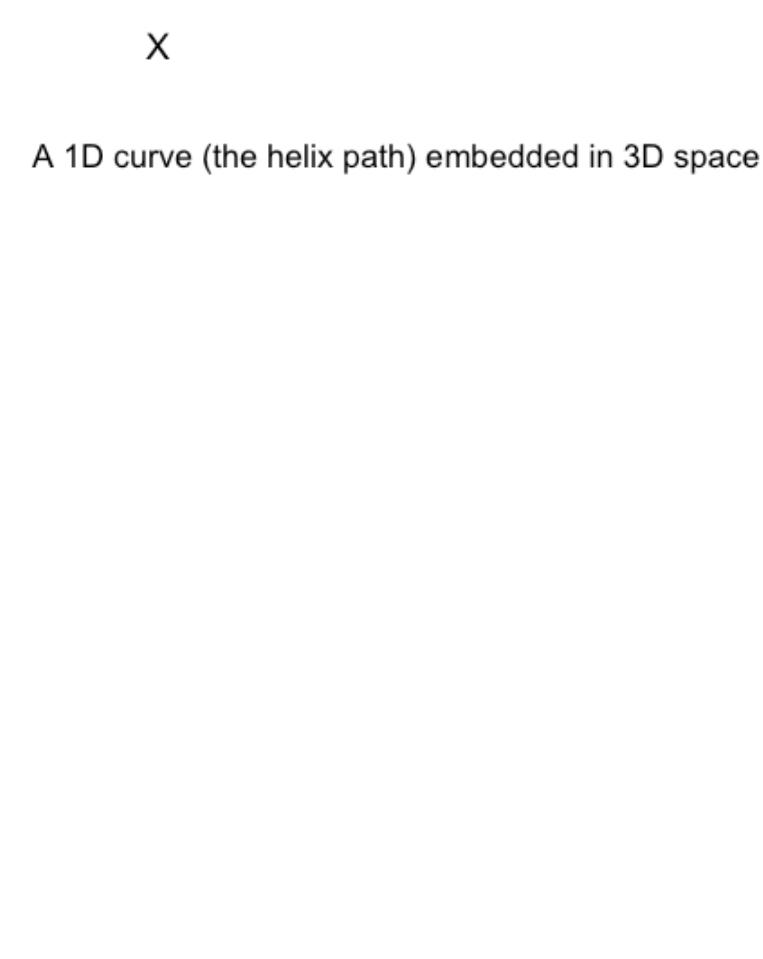
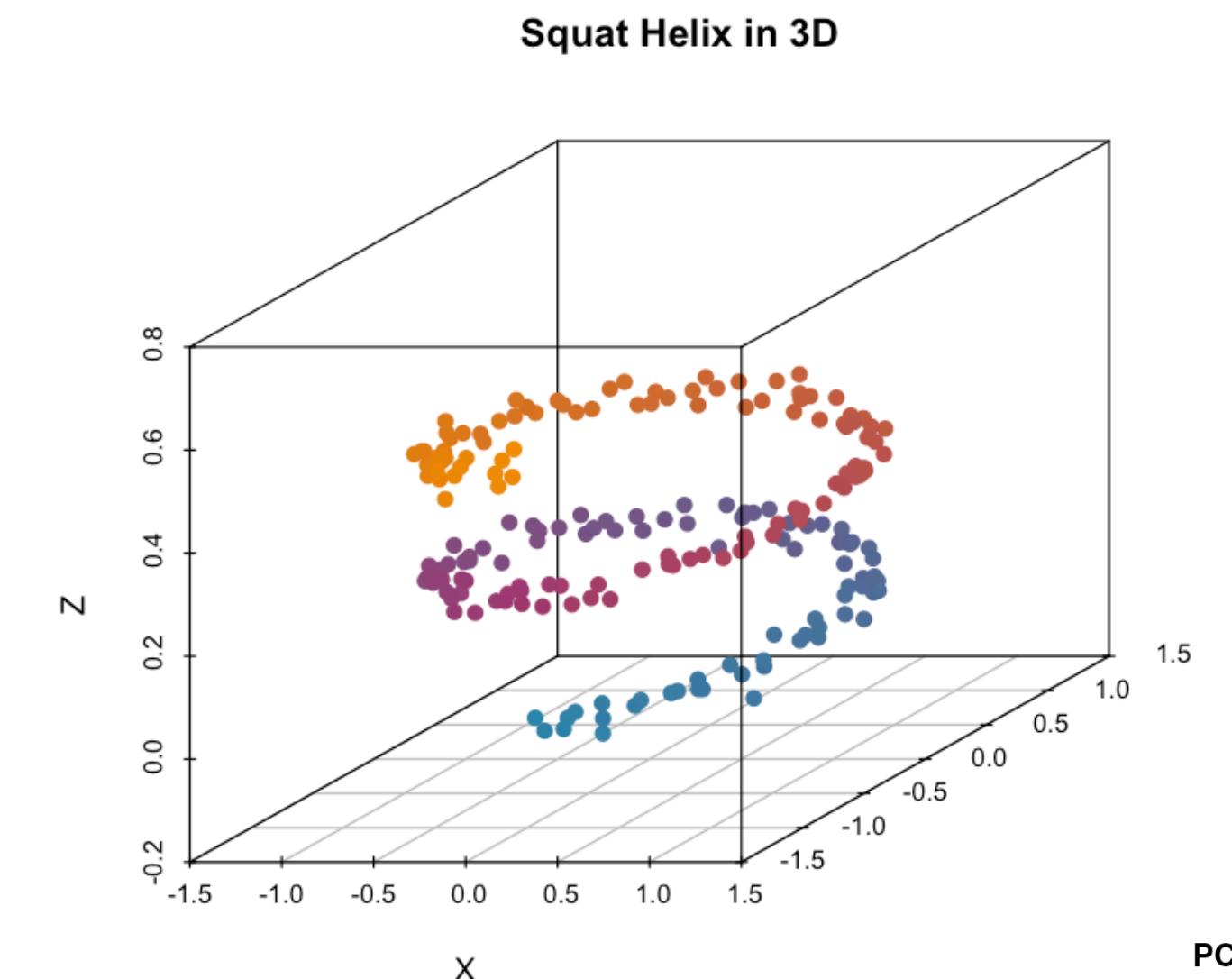


A 1D curve (the helix path) embedded in 3D space



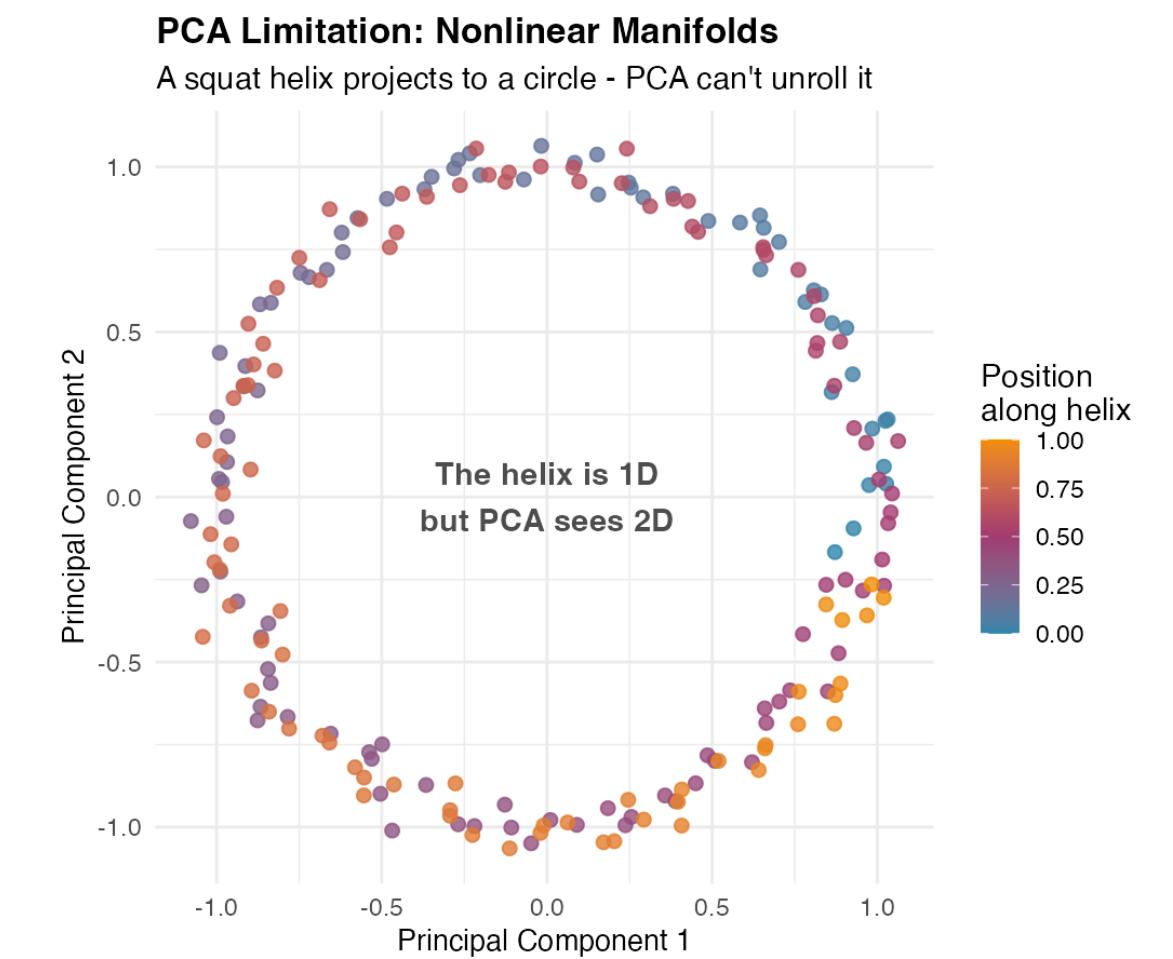
# Where PCA Fails

- PCA assumes the structure that "matters" is a **linear manifold**
  - This is often **not the case!**



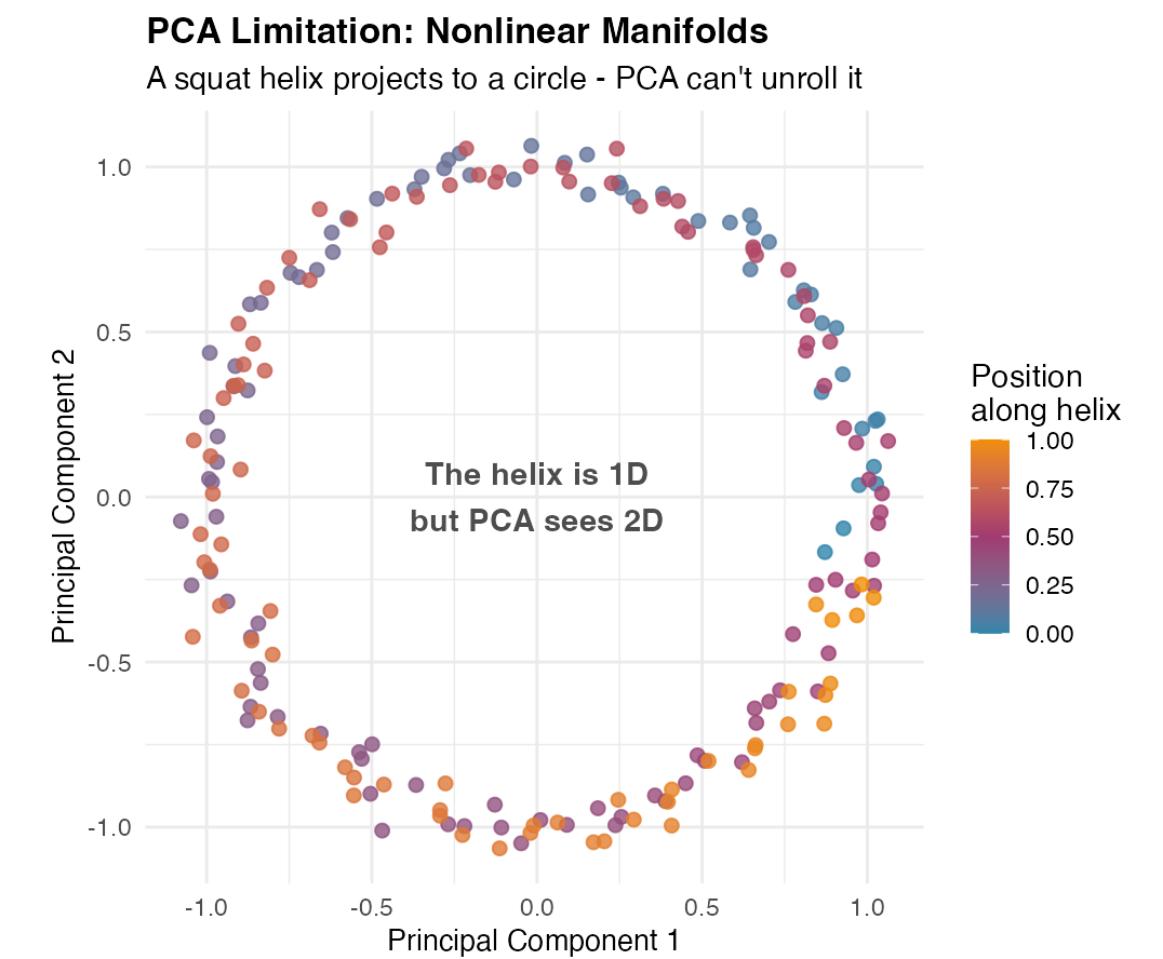
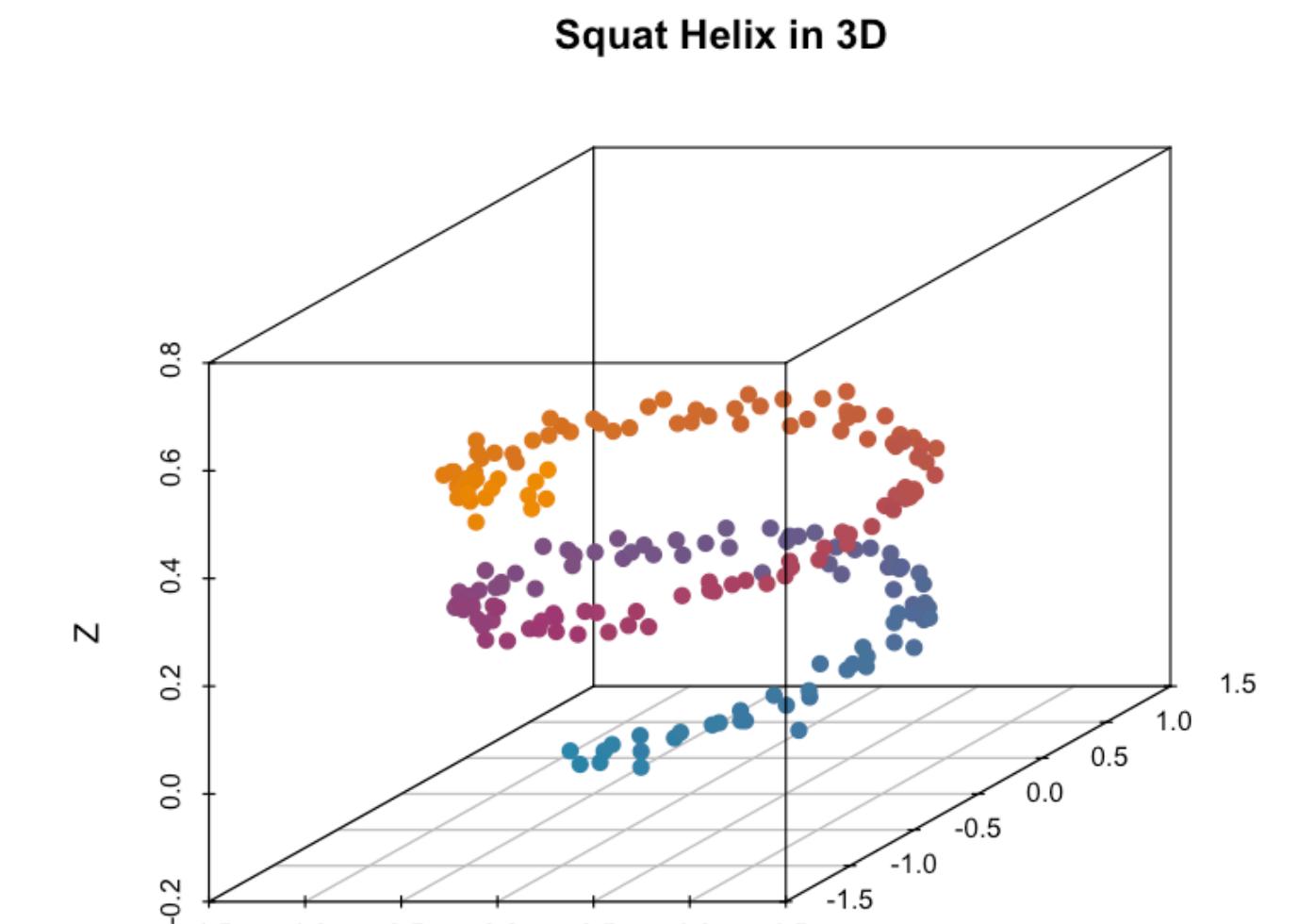
# Where PCA Fails

- PCA assumes the structure that "matters" is a **linear manifold**
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- "Squat helix": Z-dimension encodes **little variation**, but **most of the information!**
  - Technically a 1D manifold ("ascending" along the helix matters)



# Where PCA Fails

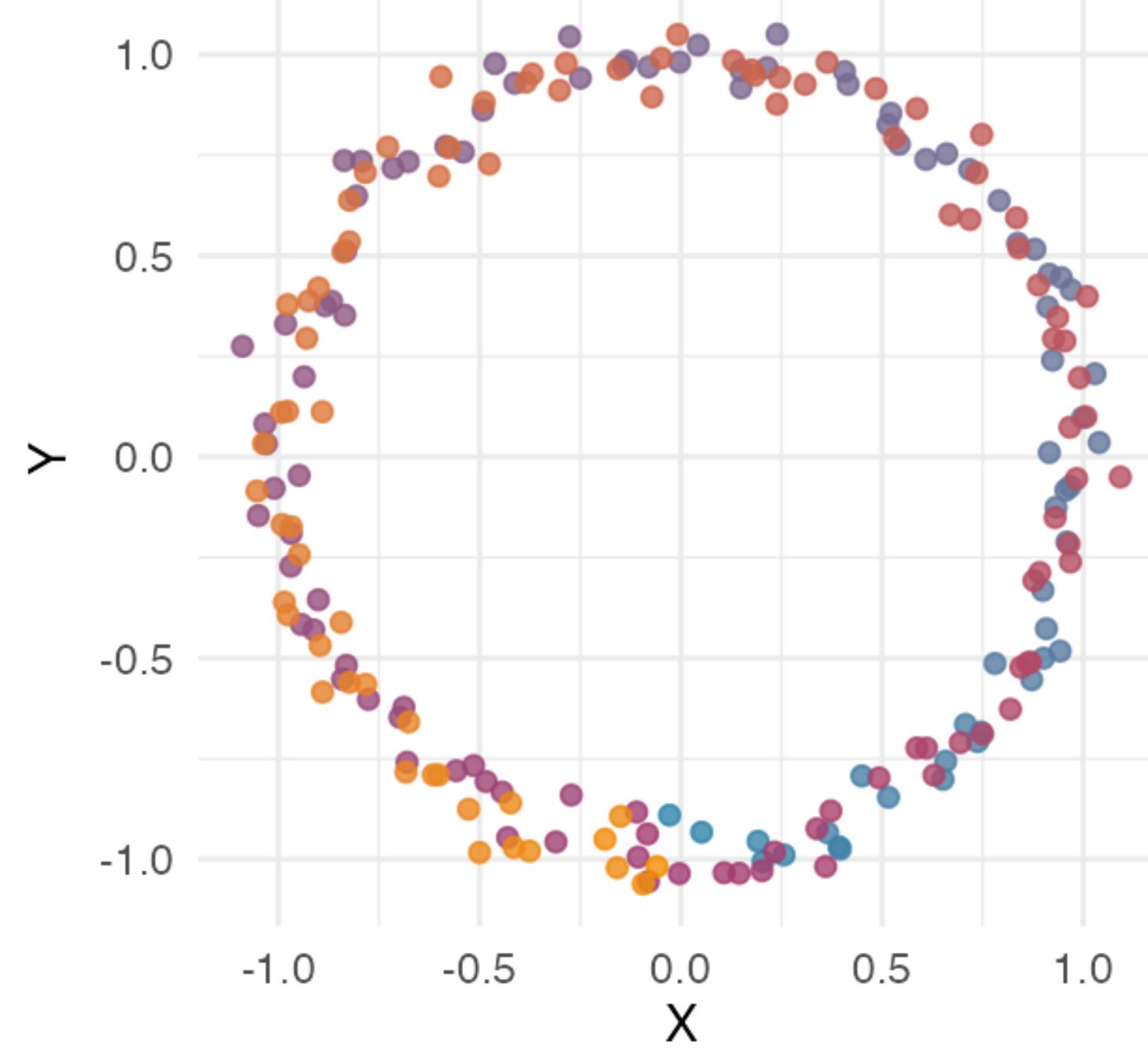
- PCA assumes the structure that "matters" is a **linear manifold**
  - This is often **not the case!**
- "Squat helix": Z-dimension encodes **little variation**, but **most of the information!**
  - Technically a 1D manifold ("ascending" along the helix matters)
- PCA will **flatten** this to a **circle**
  - **Erases** the helix information!



# Helix Example

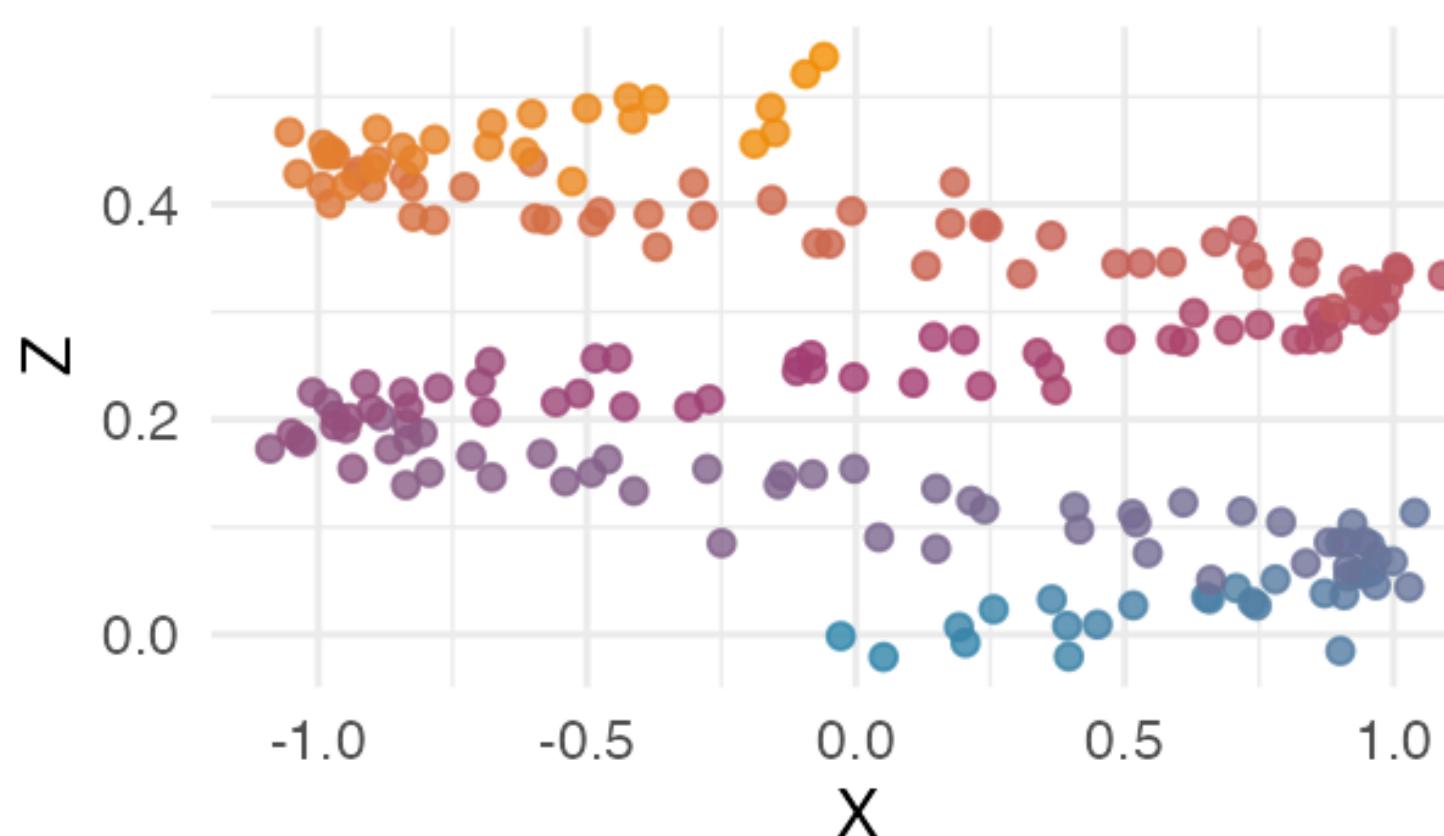
**Top-Down View (X-Y)**

Looks circular from above



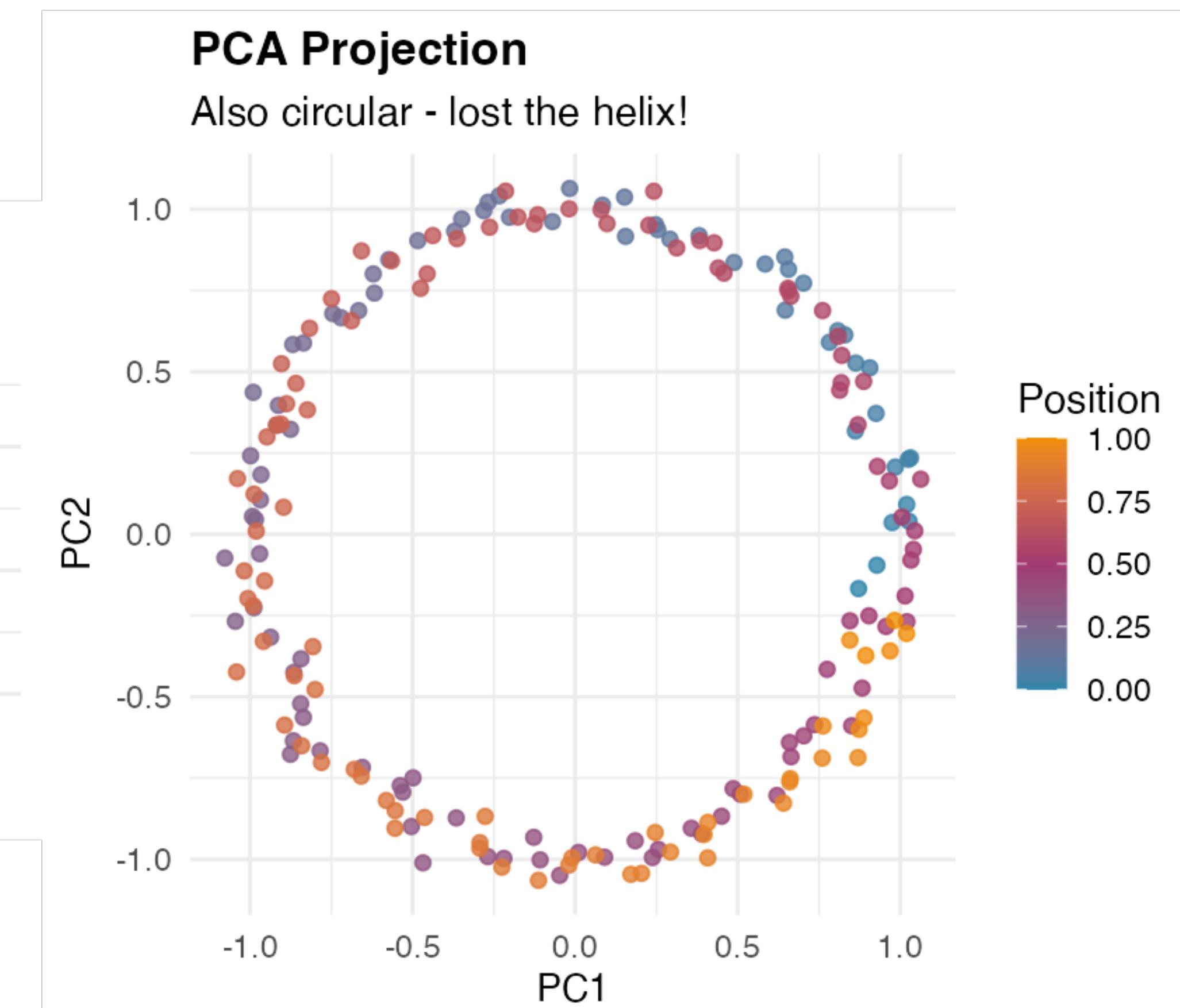
**Side View (X-Z)**

The helix structure is visible



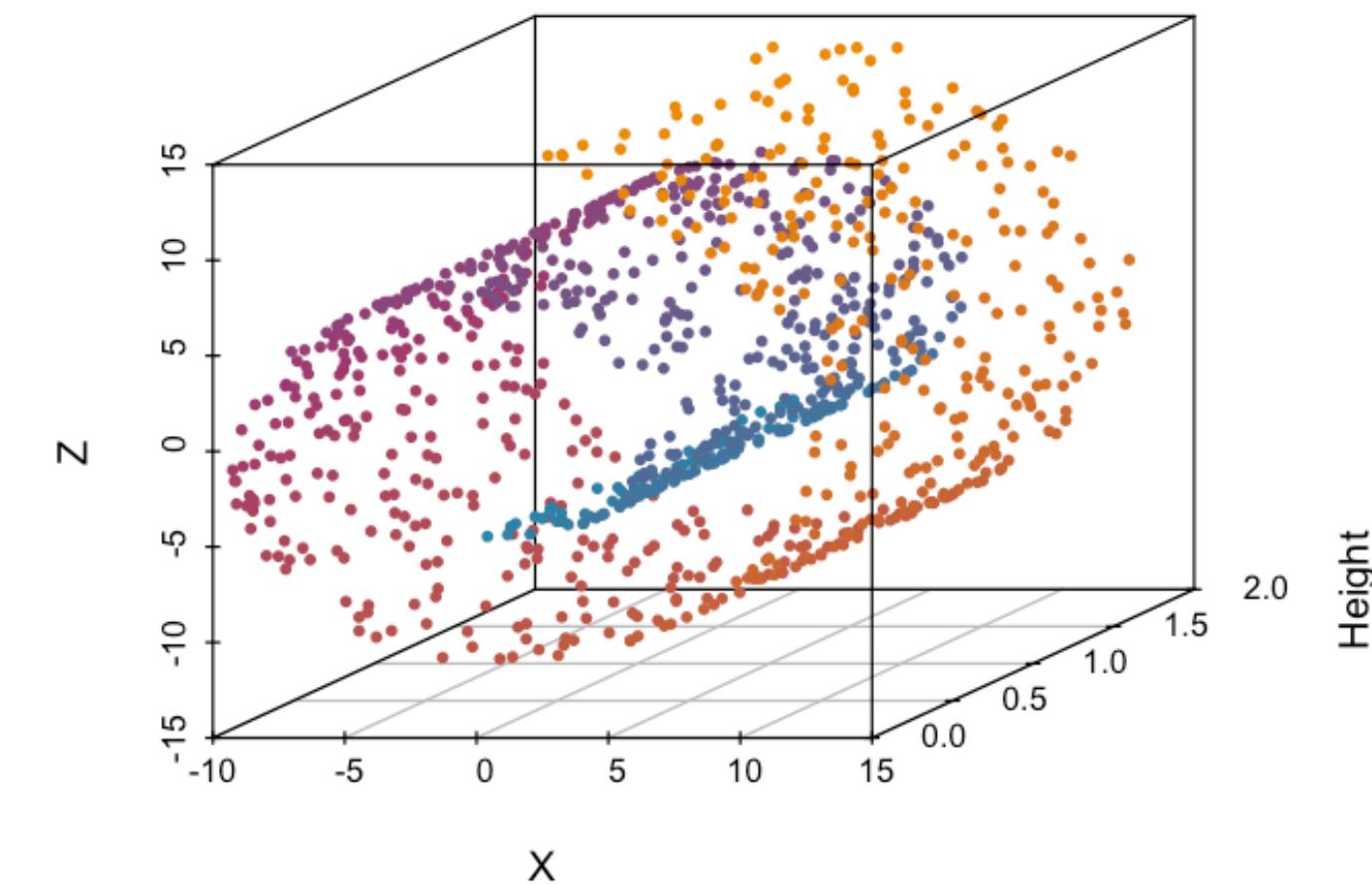
**PCA Projection**

Also circular - lost the helix!



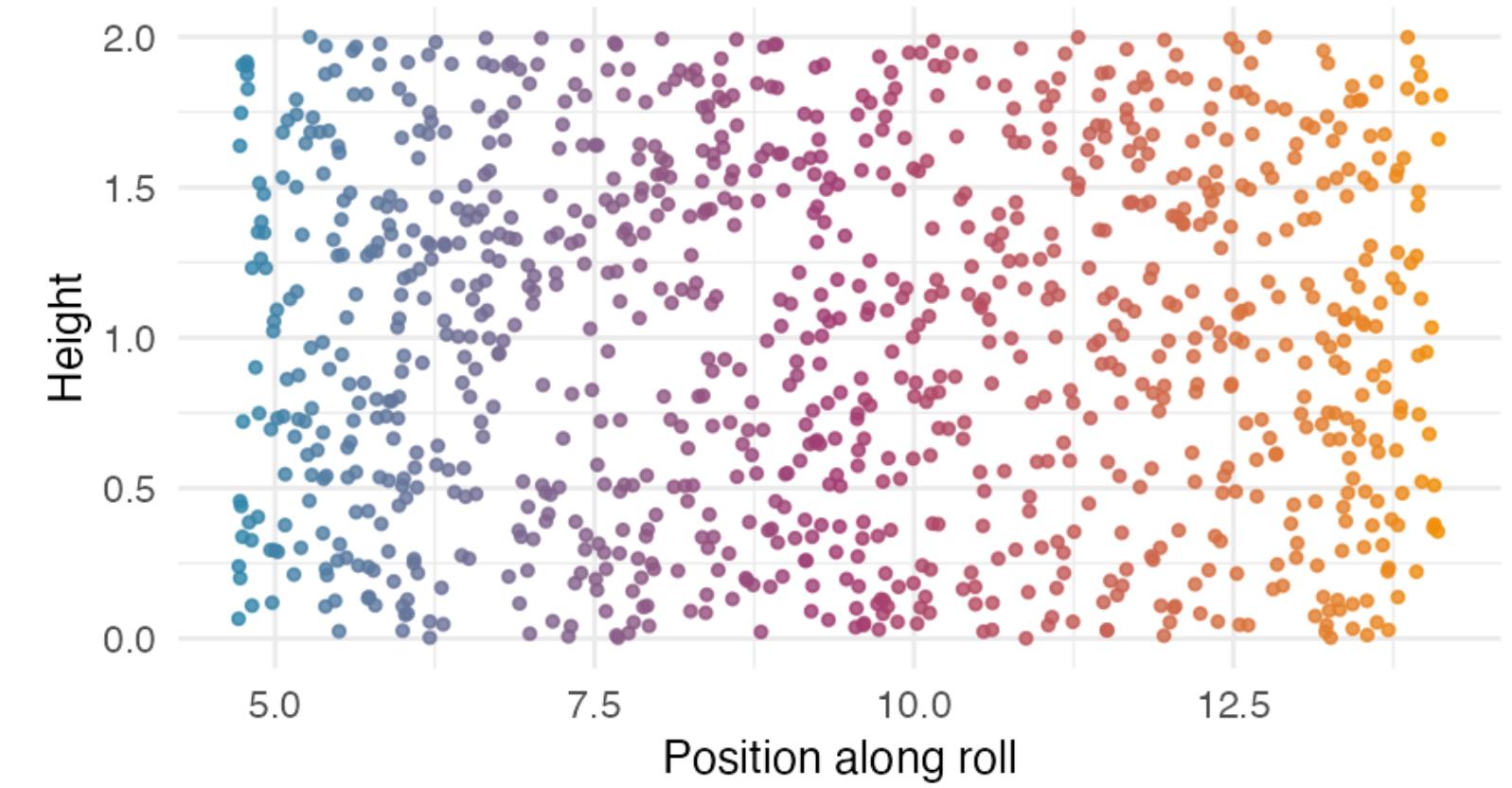
# Confound: Non-Linear Manifolds

Rolled Up: A 2D Surface in 3D



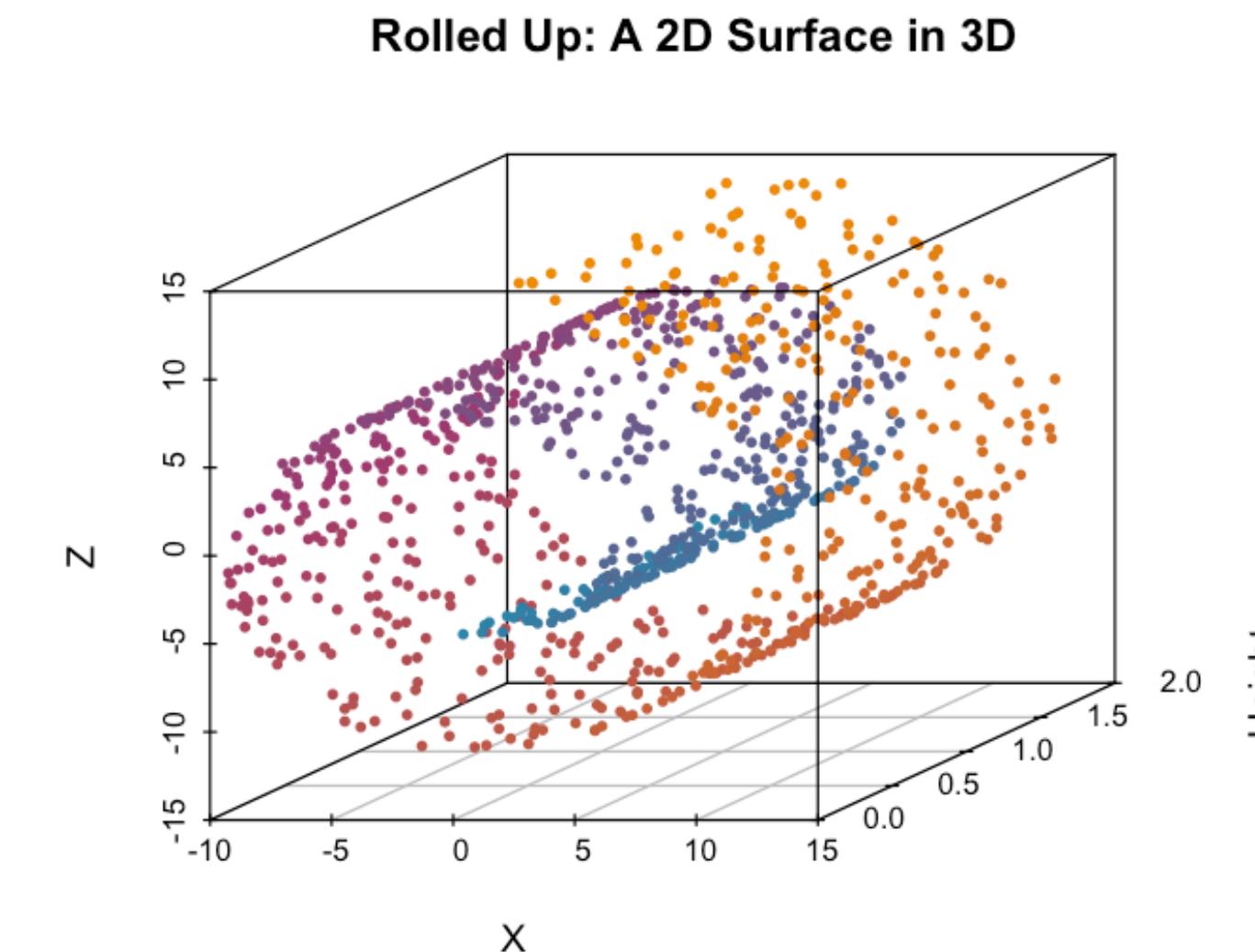
Unrolled: The True 2D Structure

What the data 'really' looks like

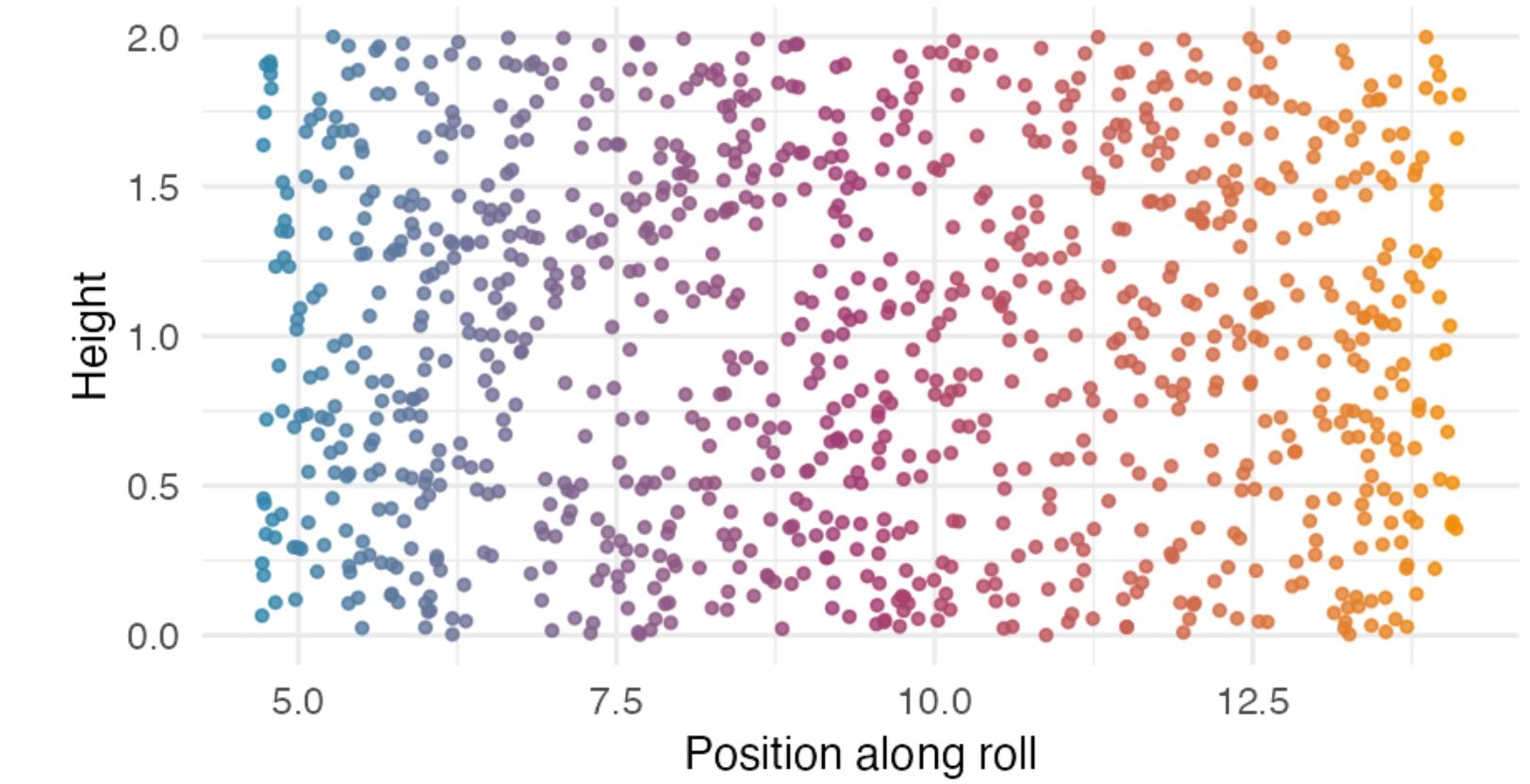


# Confound: Non-Linear Manifolds

- PCA fails when the "actual" manifold is **non-linear**
  - I.e. the **inductive bias** hurts us!

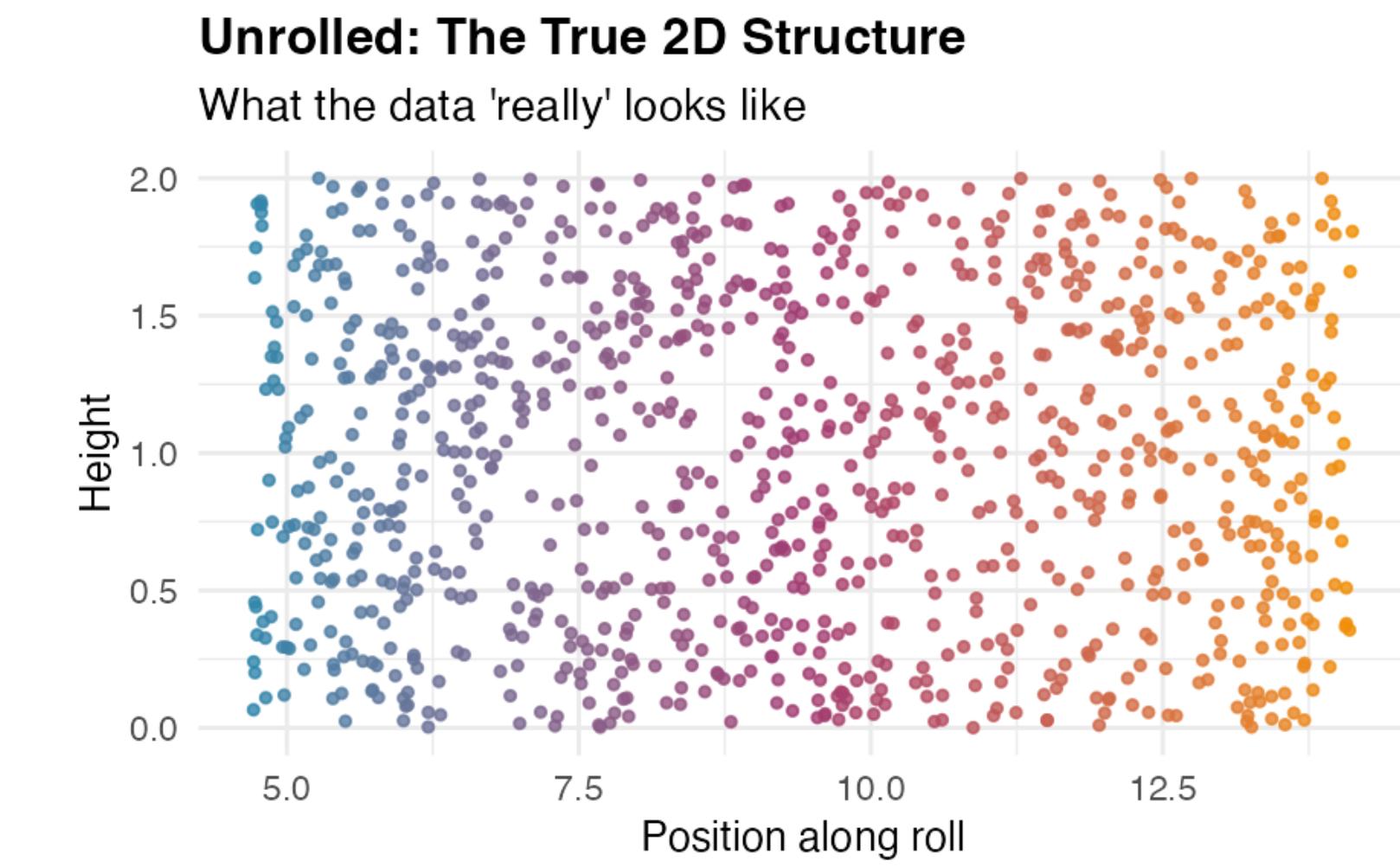
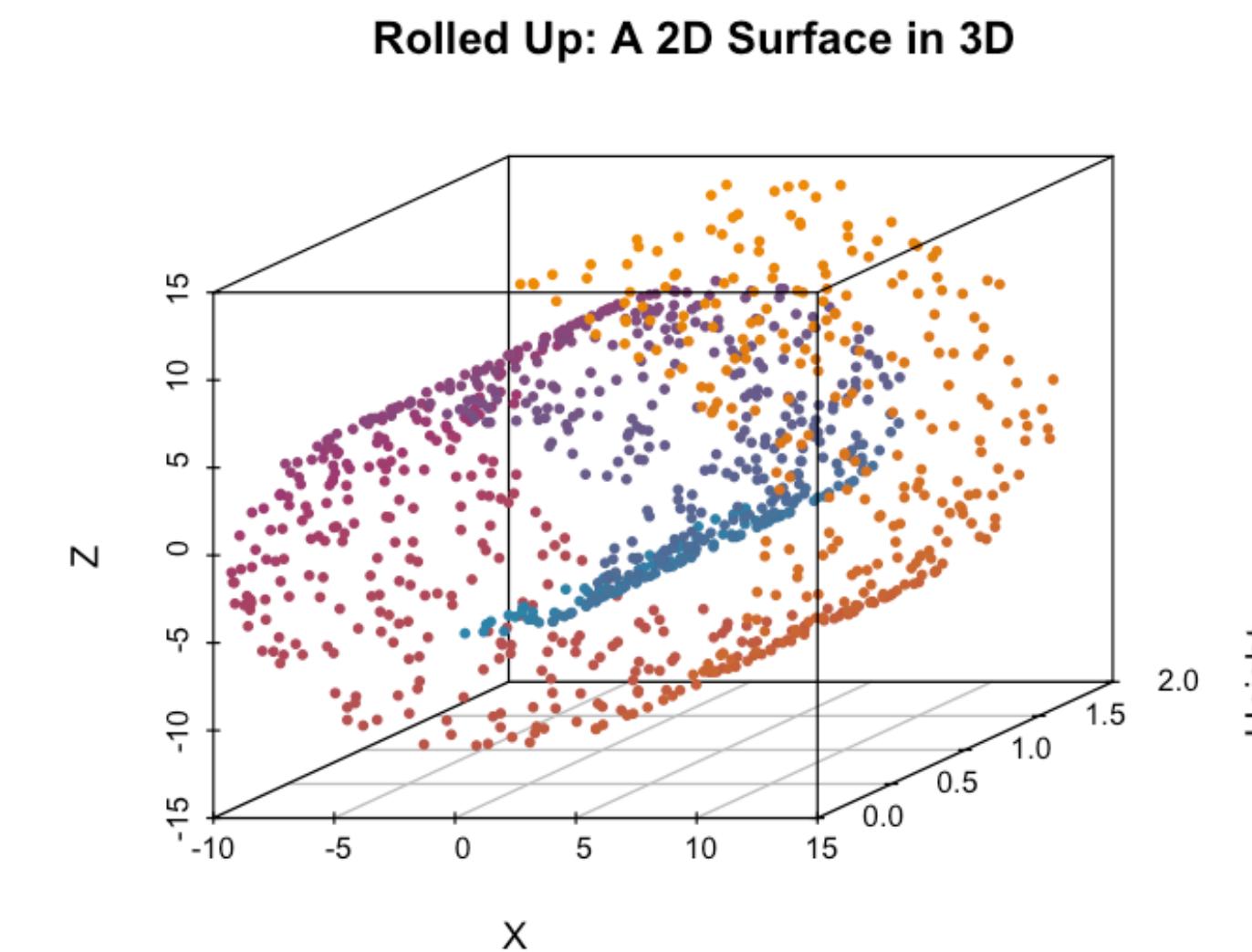


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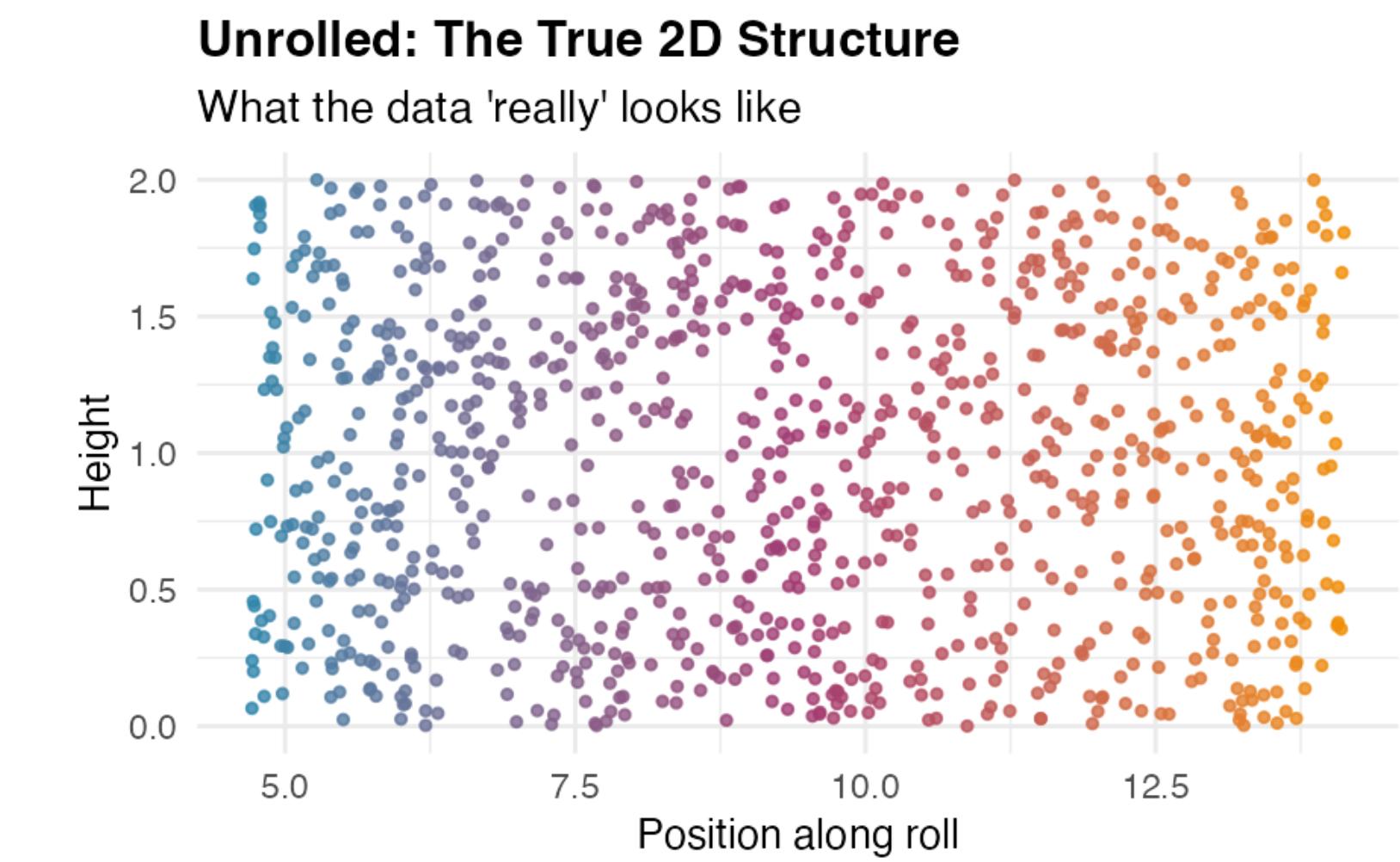
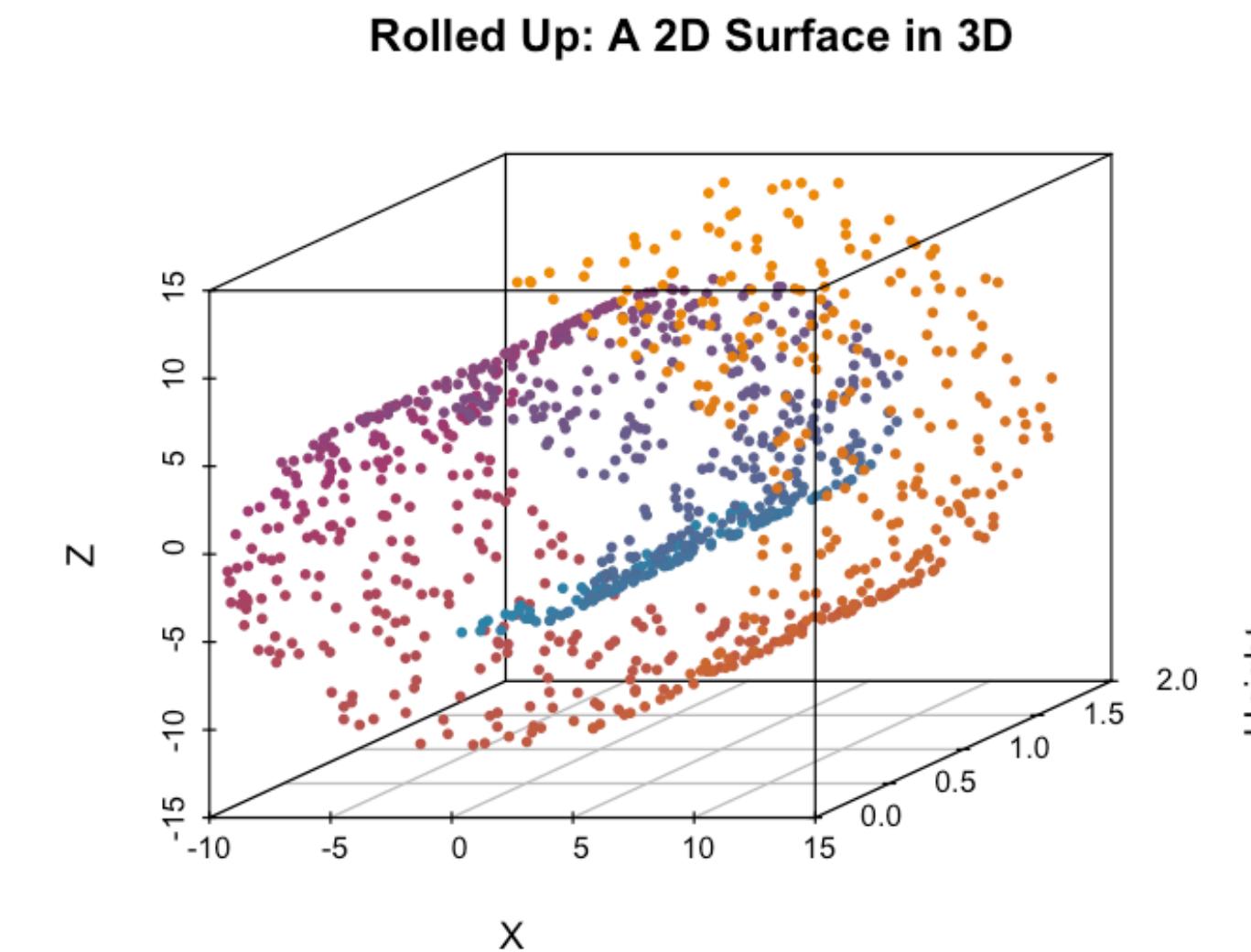
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  - E.g. "rolling up" a 2D surface

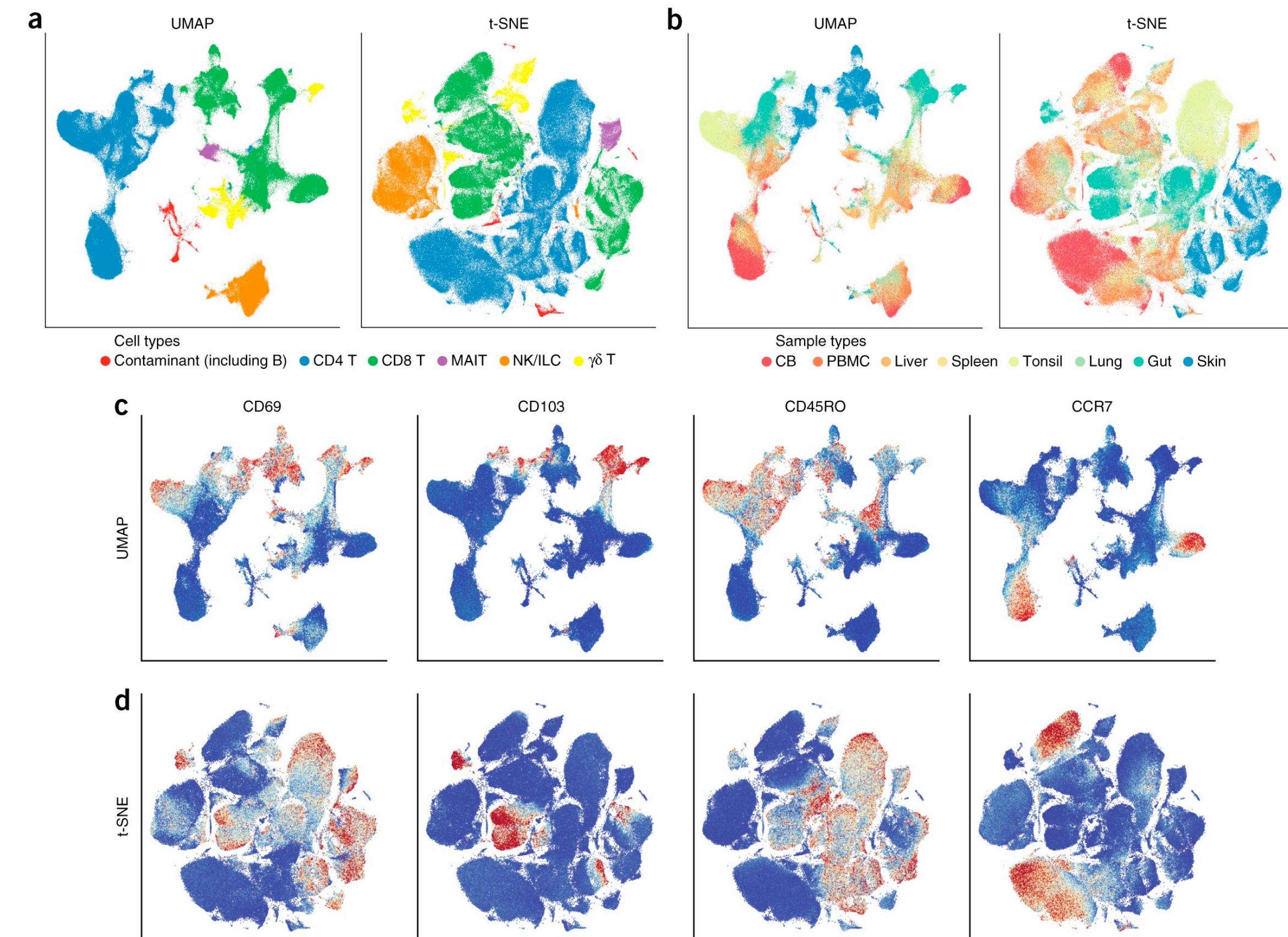


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  - I.e. the **inductive bias** hurts us!
- Manifolds can have **arbitrarily complex shapes**
  - E.g. "rolling up" a 2D surface
- How do we deal with this?
  - Algorithms to **learn non-linear manifolds**

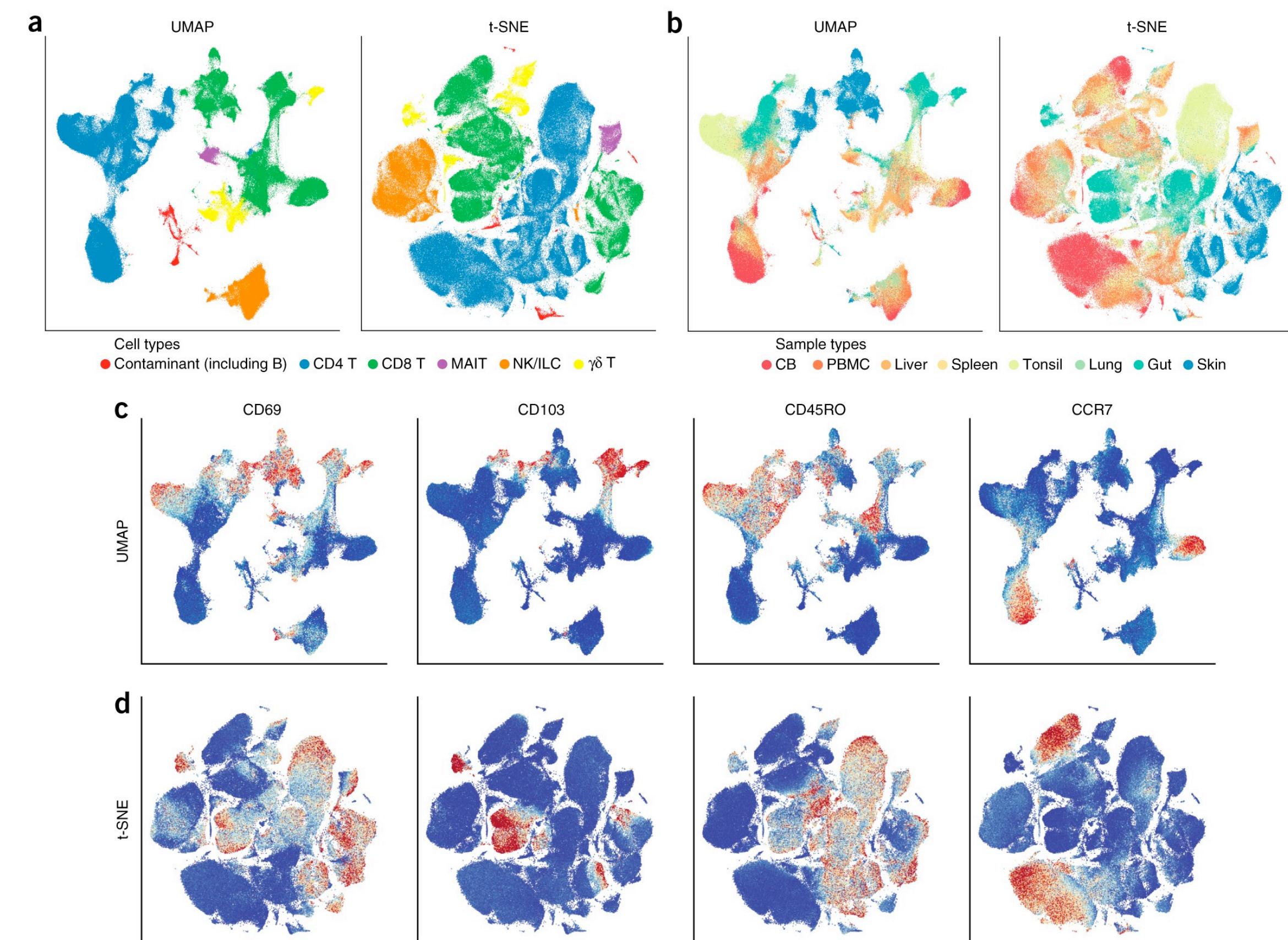


# Related Non-Linear Algorithms



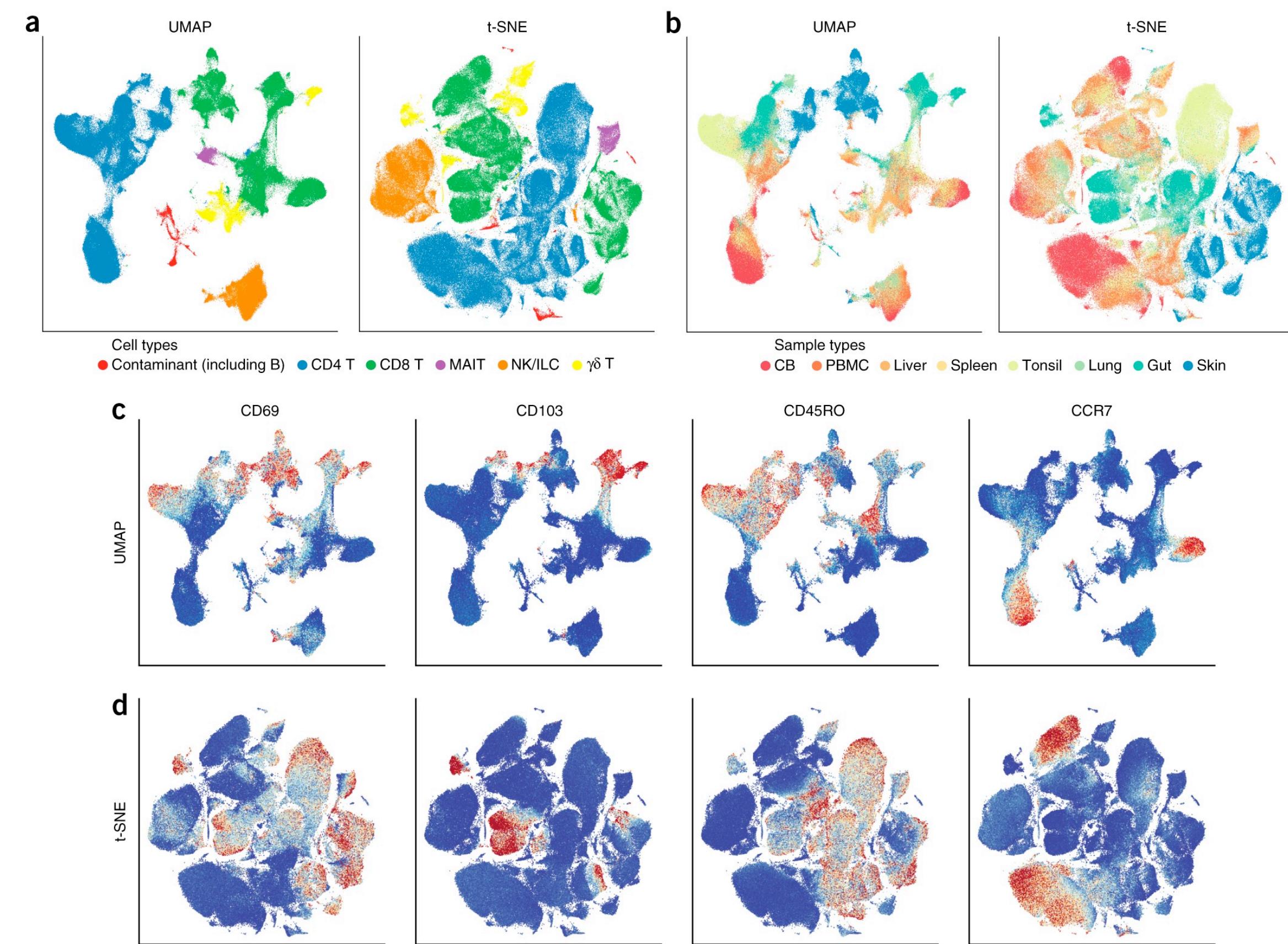
# Related Non-Linear Algorithms

- t-SNE and UMAP resemble PCA input/output but...
  - They can learn **non-linear manifolds**
  - Focus more on preserving **local structure** where PCA preserves global structure



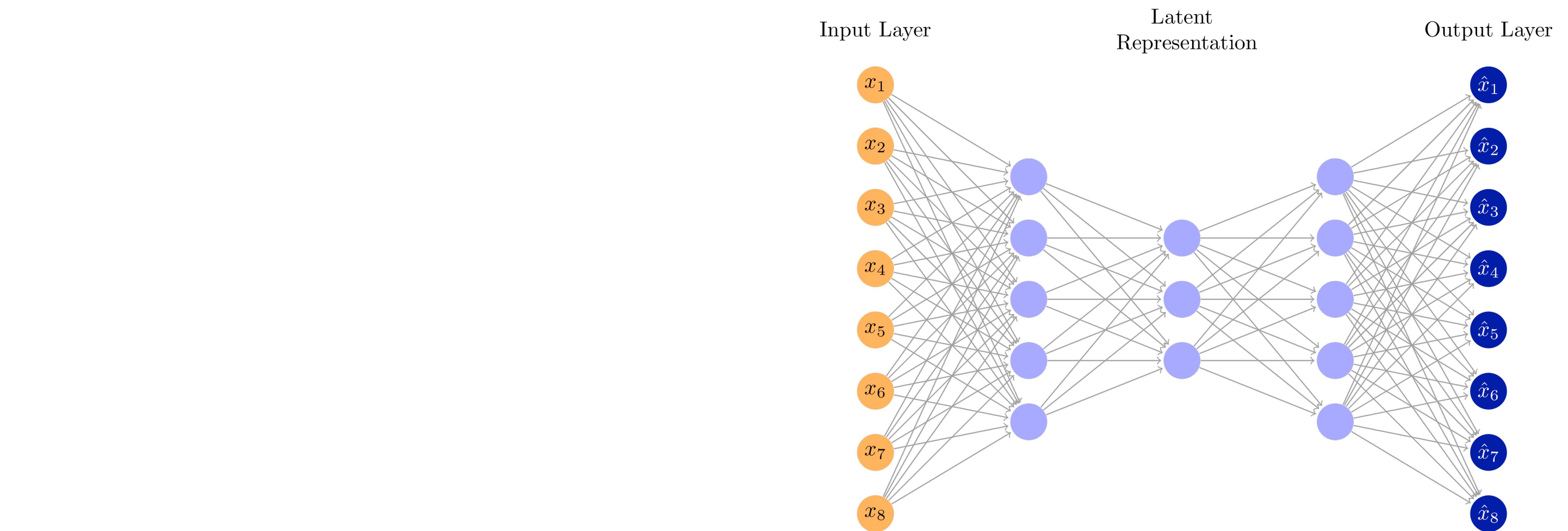
# Related Non-Linear Algorithms

- t-SNE and UMAP resemble PCA input/output but...
  - They can learn **non-linear manifolds**
  - Focus more on preserving **local structure** where PCA preserves global structure
- Caveats: mostly good for **visualization** and **exploratory analysis**
  - The distance between output clusters is **not interpretable**
  - Very sensitive to **hyper-parameters**
  - Output features tend to be **not good for downstream ML**



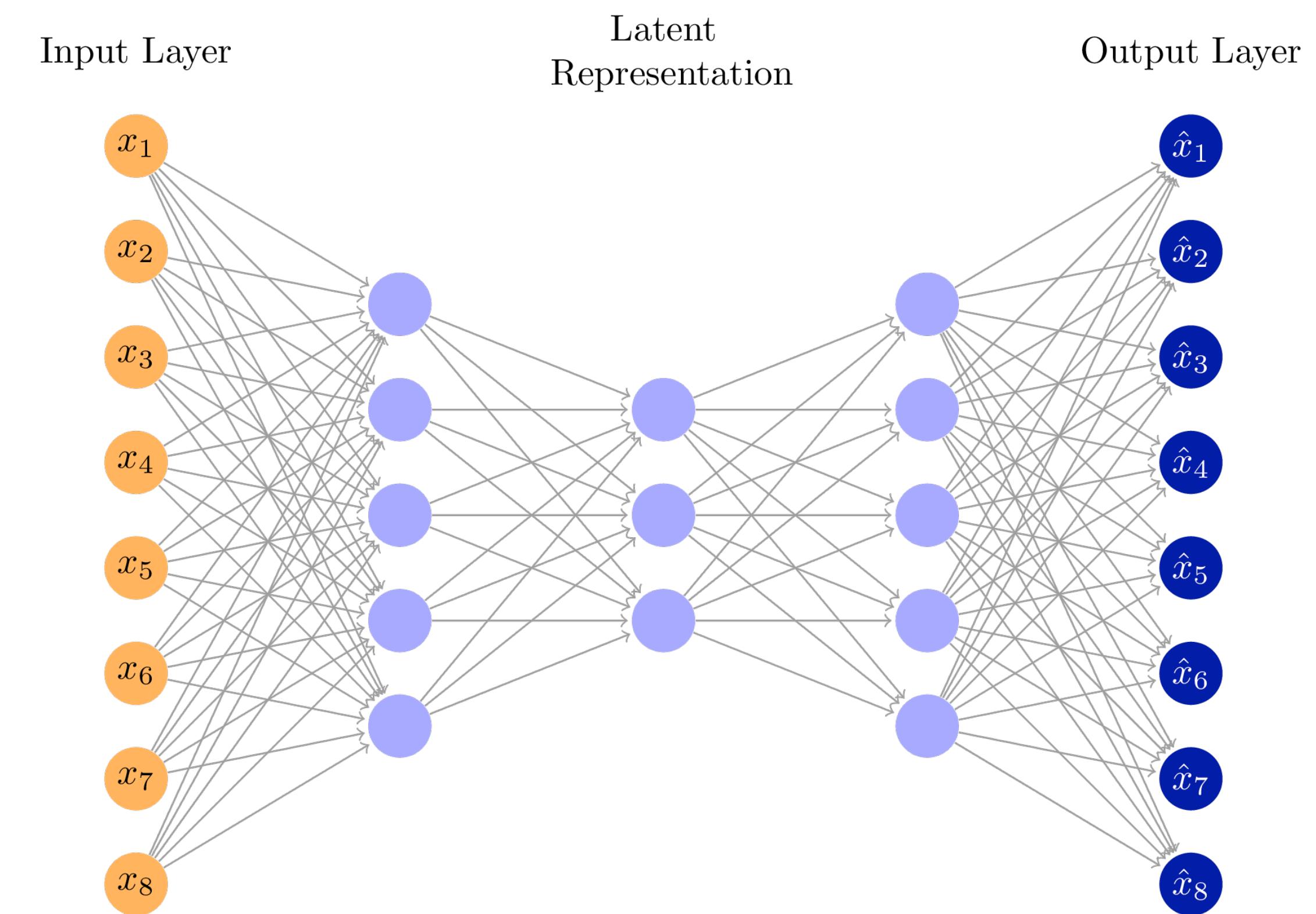
# Autoencoders

# Autoencoders: Basic Idea



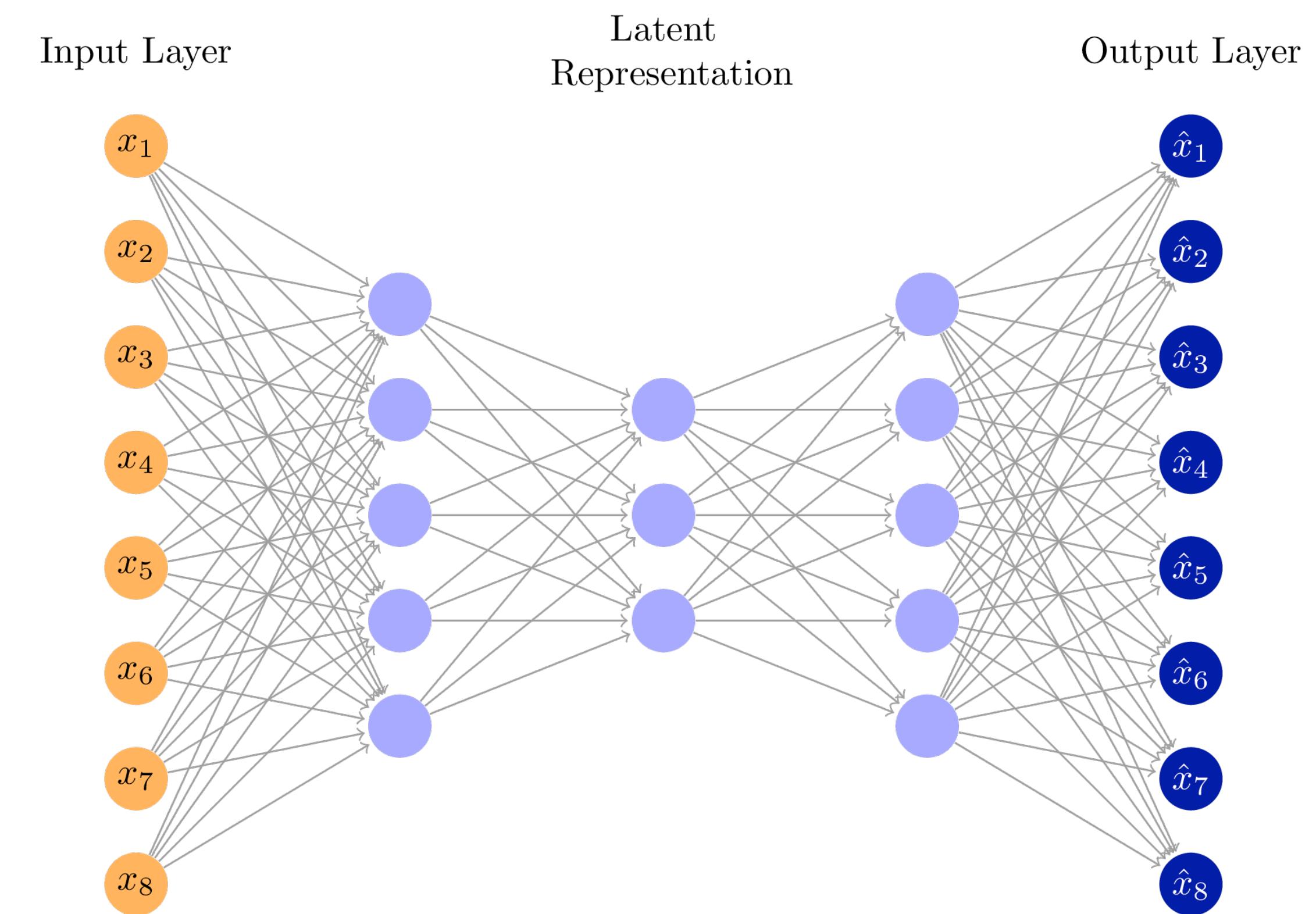
# Autoencoders: Basic Idea

- Autoencoders are **Neural Network-based** algorithms that **learn to compress data**

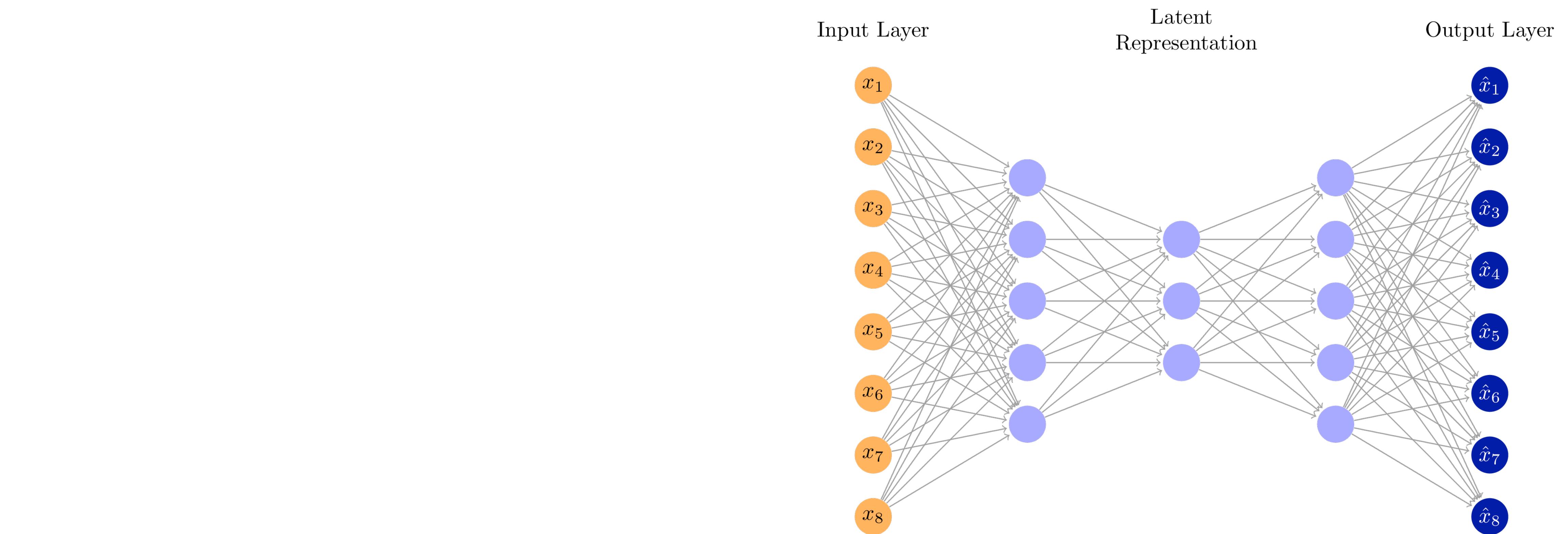


# Autoencoders: Basic Idea

- Autoencoders are **Neural Network-based** algorithms that **learn to compress data**
- Learn a low-dimensional **bottleneck representation**, which can be used to **reconstruct the input**
  - This forces dimensionality reduction with **minimal loss of information**

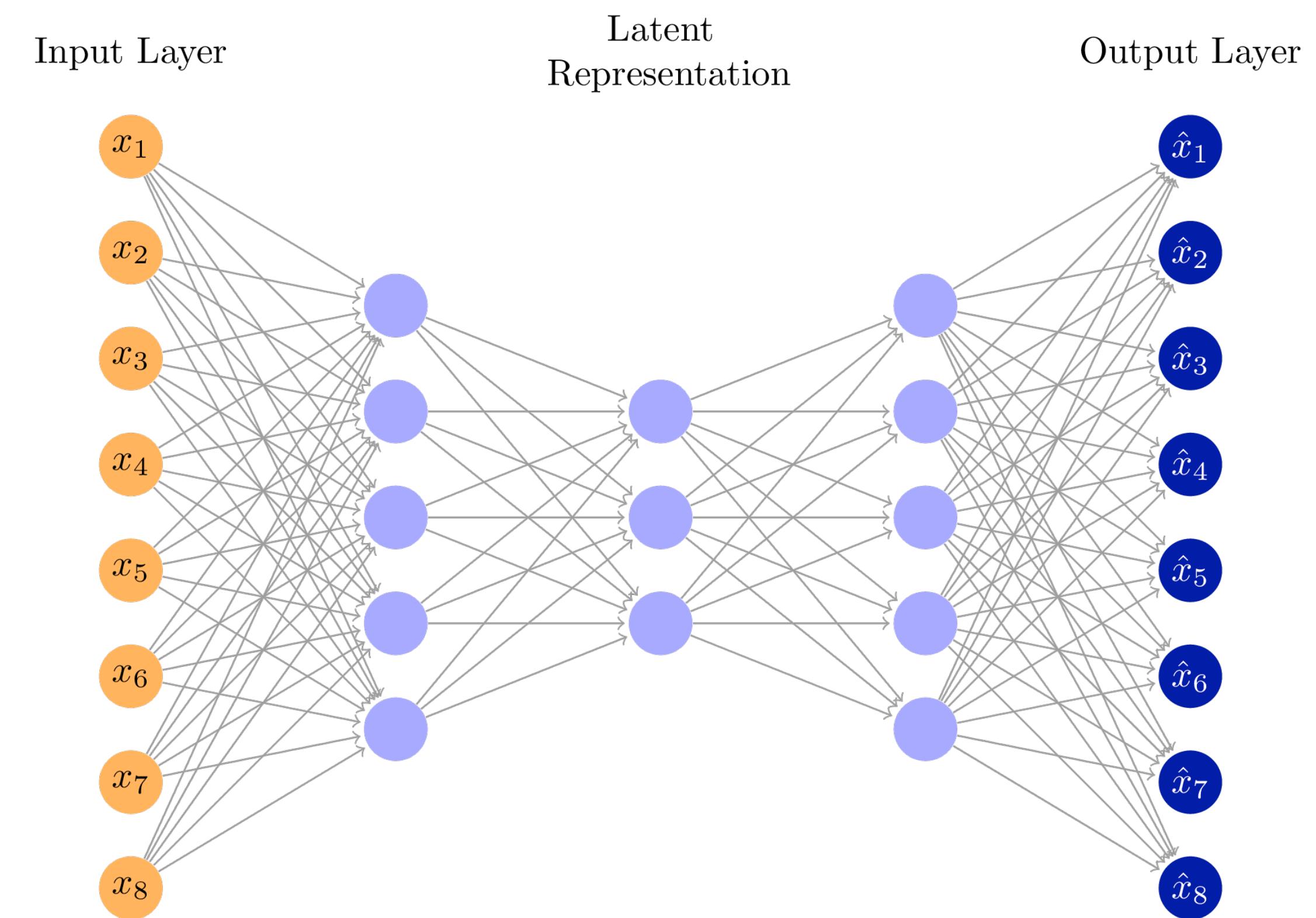


# Why does this work?



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- Can learn non-linear manifolds because **NNs apply successive non-linear transformations**
- Can think of it as NN learning to "unravel" or "untangle" the complex non-linear manifold



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- Can learn non-linear manifolds because **NNs apply successive non-linear transformations**
- Can think of it as NN learning to "unravel" or "untangle" the complex non-linear manifold
- High-dimensional representation is **simplified** to what we hope is its "intrinsic" dimension

