

# PCA Review, Probabilistic PCA, GPLVMs

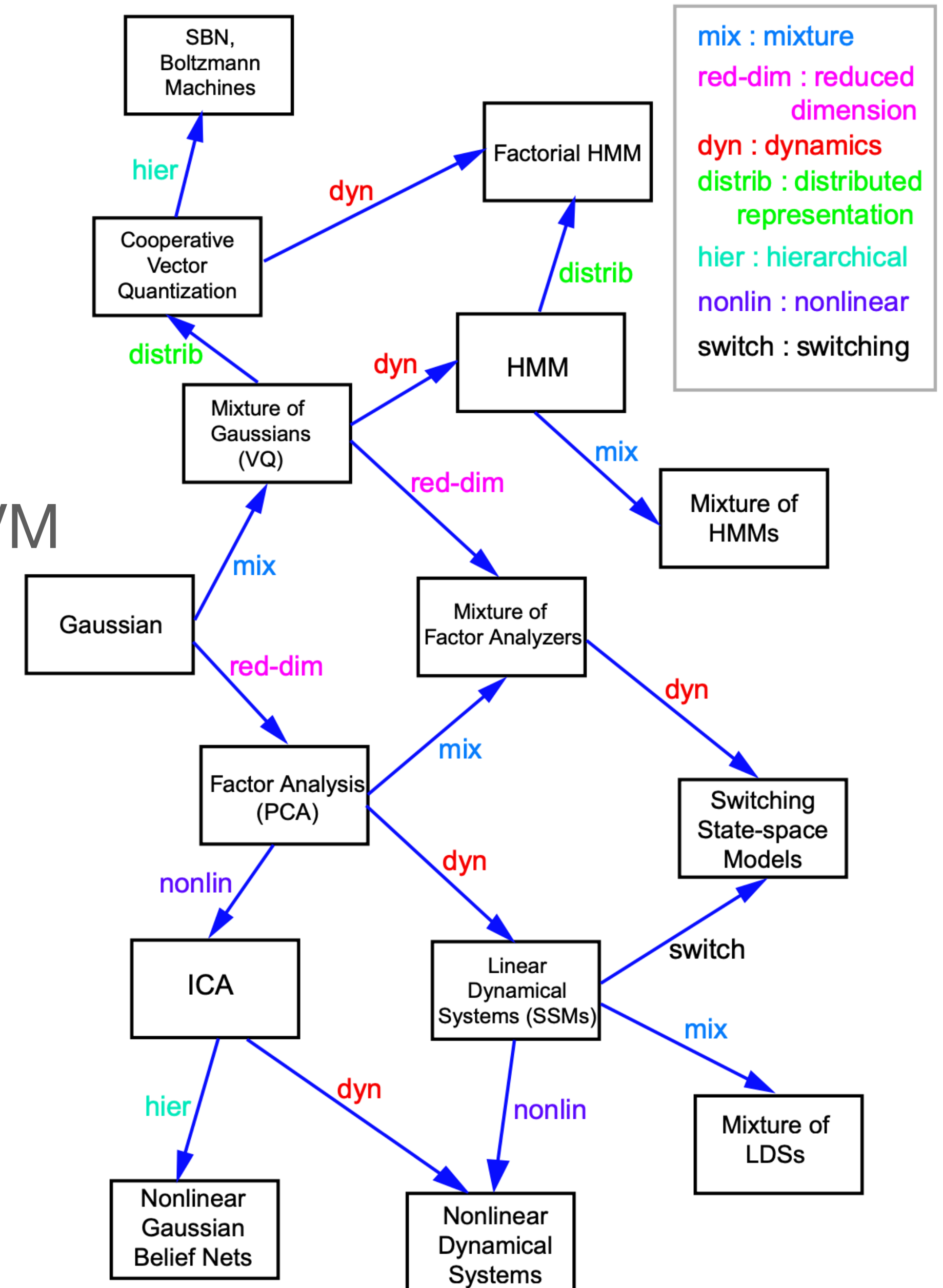
# Housekeeping

- + Quiz 1 results are available on Wattle, 9.5/10 average!
- + Class survey is running. Also feedback [direct (emails) or indirect (class reps)] are appreciated.
- + Assignment 1 due this Friday [midnight]. 5% penalty for 5 mins - 24 hrs late, 100% penalty after.
- + I will be on leave the following two weeks so emails will be slow
- + Prof Jing Jiang will take over after the break.

# Overview

1. Motivation
2. PCA review
3. Linear, Gaussian latent variable models and GPLVM

Reading: Bishop 12.1, 12.2, 12.4.2



# Motivation

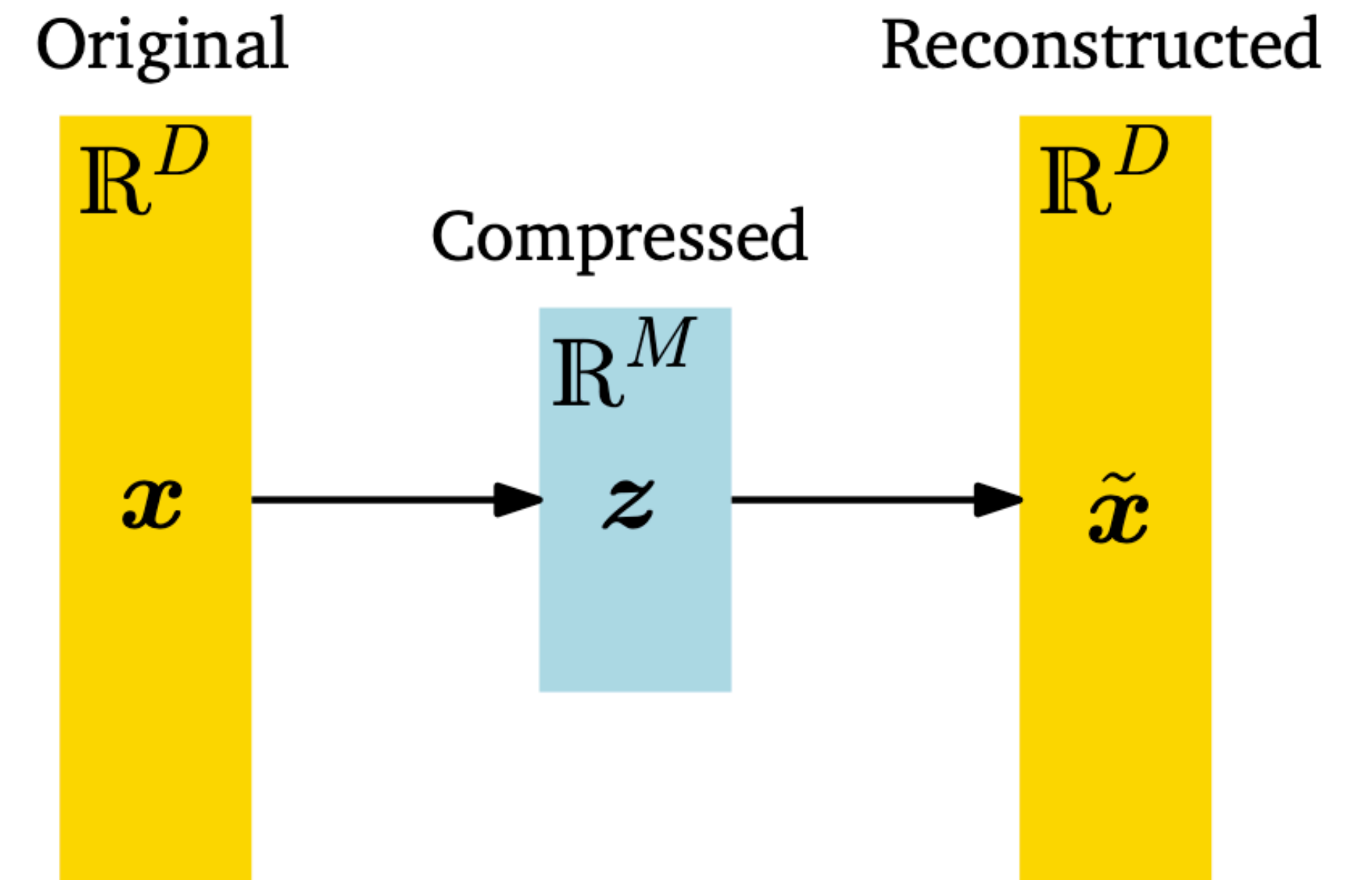
# Motivation

## Dimensionality reduction as data compression

Find lower-dimensional data without losing much information

$$M < D$$

$\mathbf{z}$  captures desirable variations in  $\mathbf{x}$   
Reconstructed data is similar to  $\mathbf{x}$



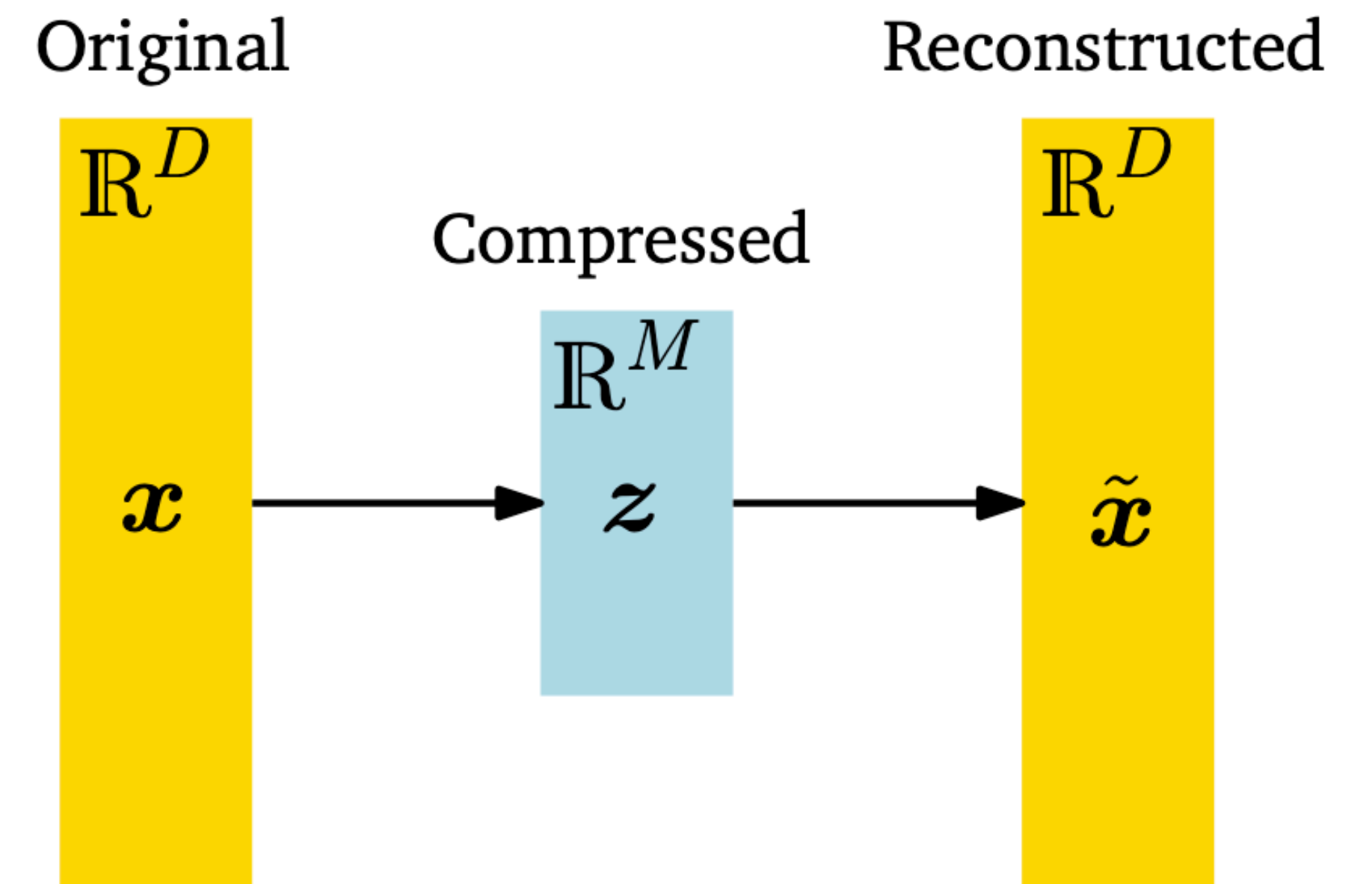
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## Why?

- + Data may have low *intrinsic* dimensionality [think about data living on a line in high dimensions]
- + visualisation / exploratory data analysis [e.g. compress 100-D data down to 2D to visualise patterns]
- + Using low dimensional data for learning [e.g. train a classifier using compressed data]

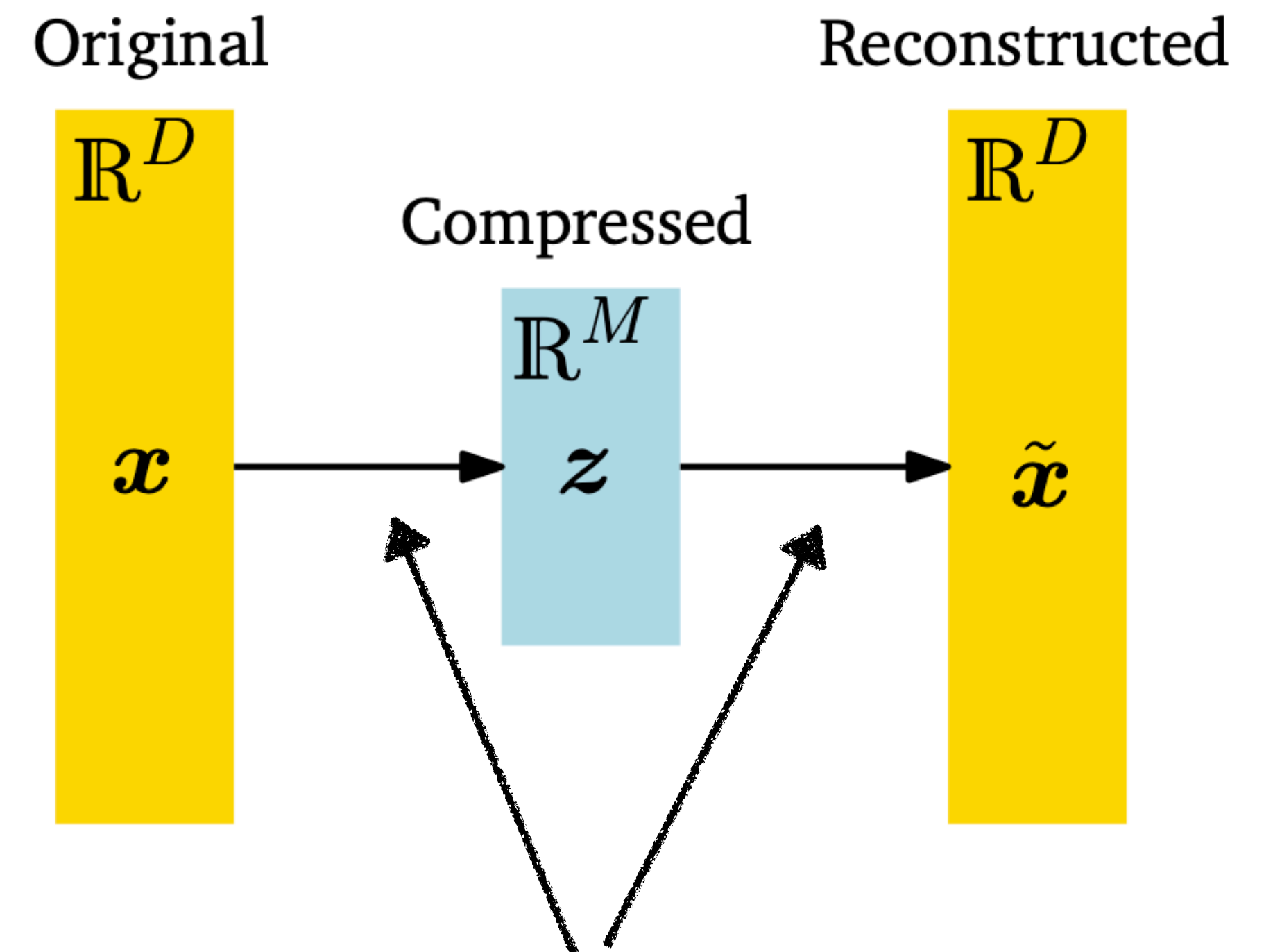
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## Why?

**Key question: how to construct these mappings?**

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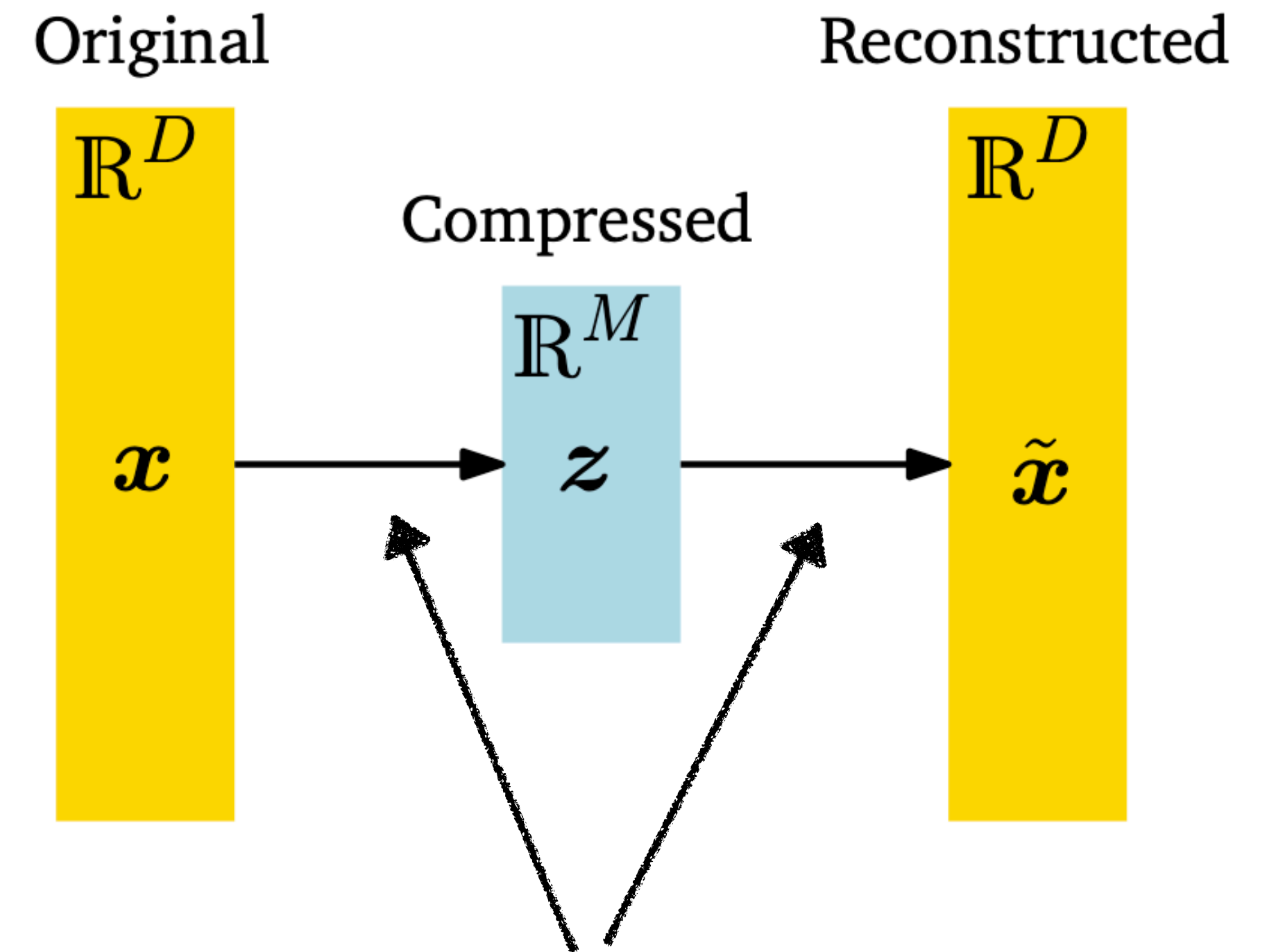
# Motivation - example

## Dimensionality reduction as data compression

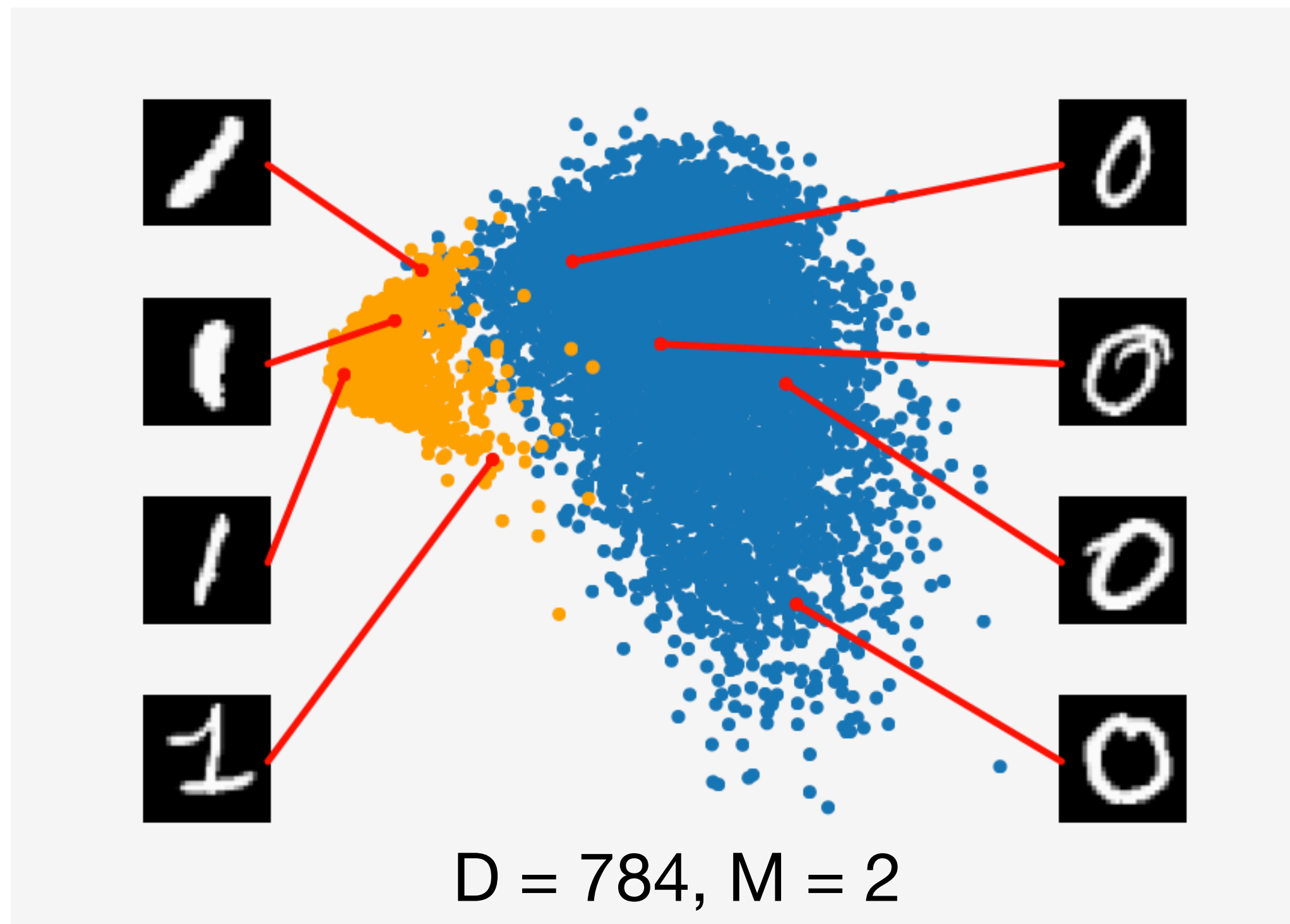
Find lower-dimensional data without losing much information

$$M < D$$

$z$  captures desirable variations in  $x$   
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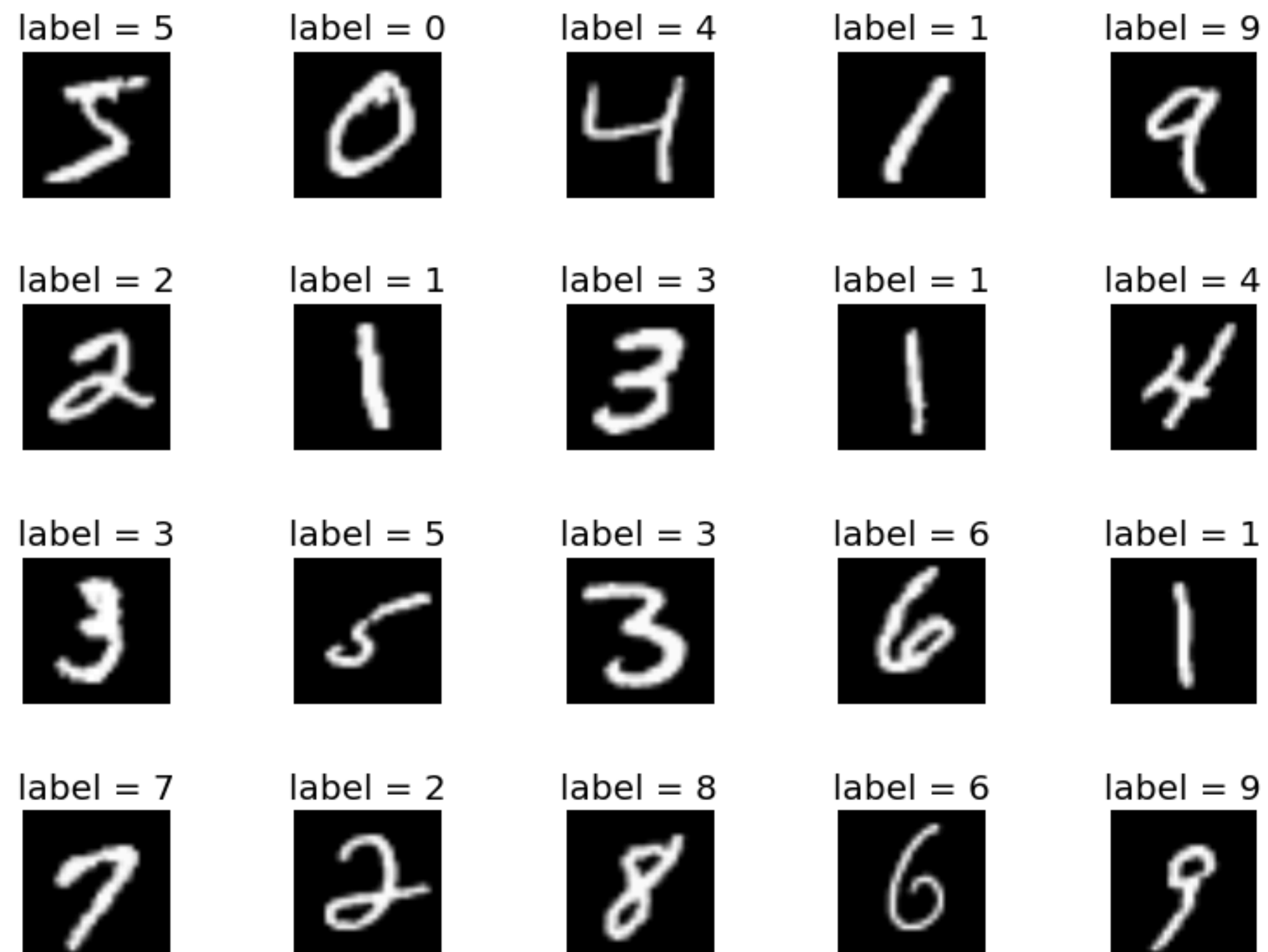
Key question: how to construct these mappings?



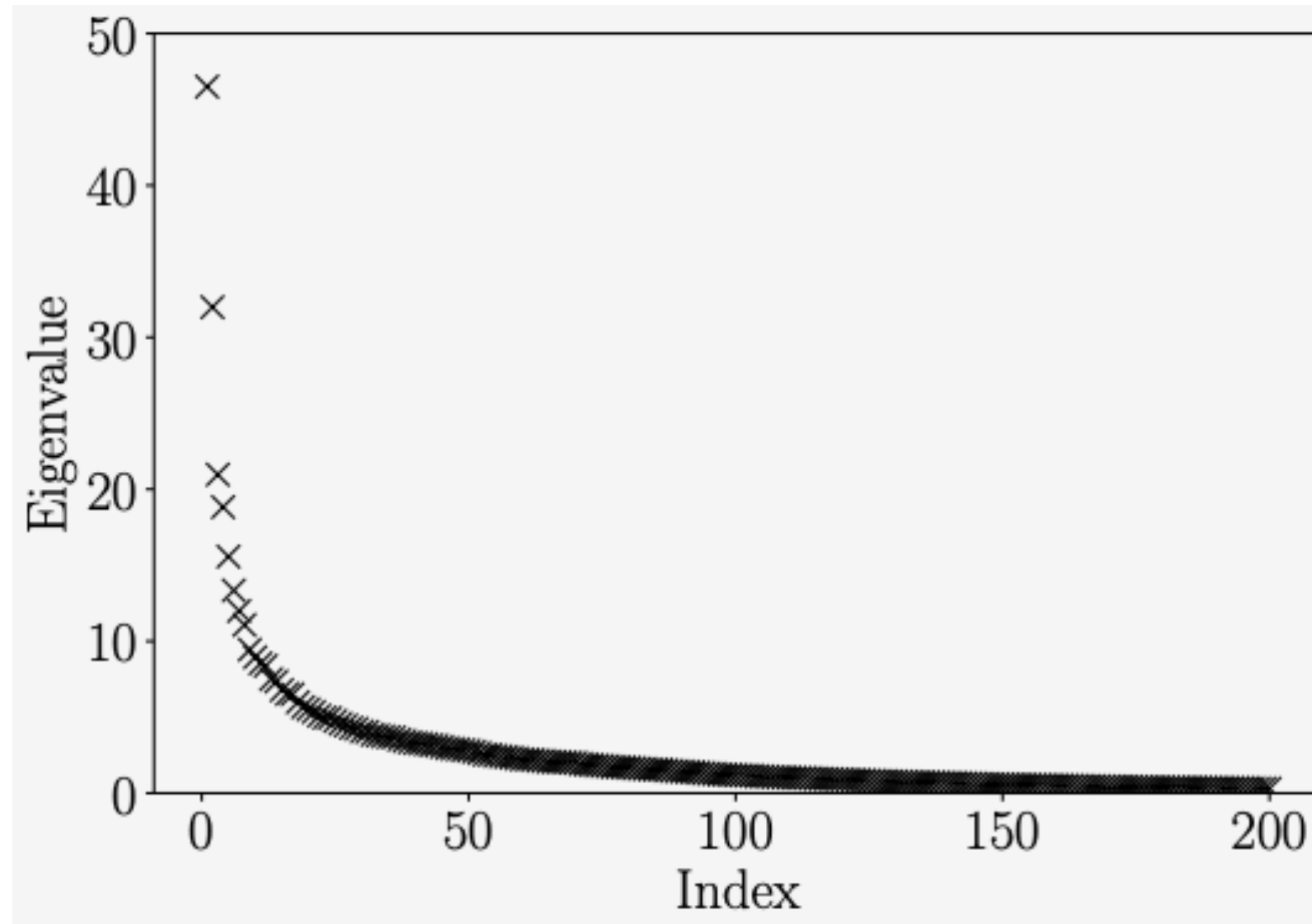


# Example - dataset

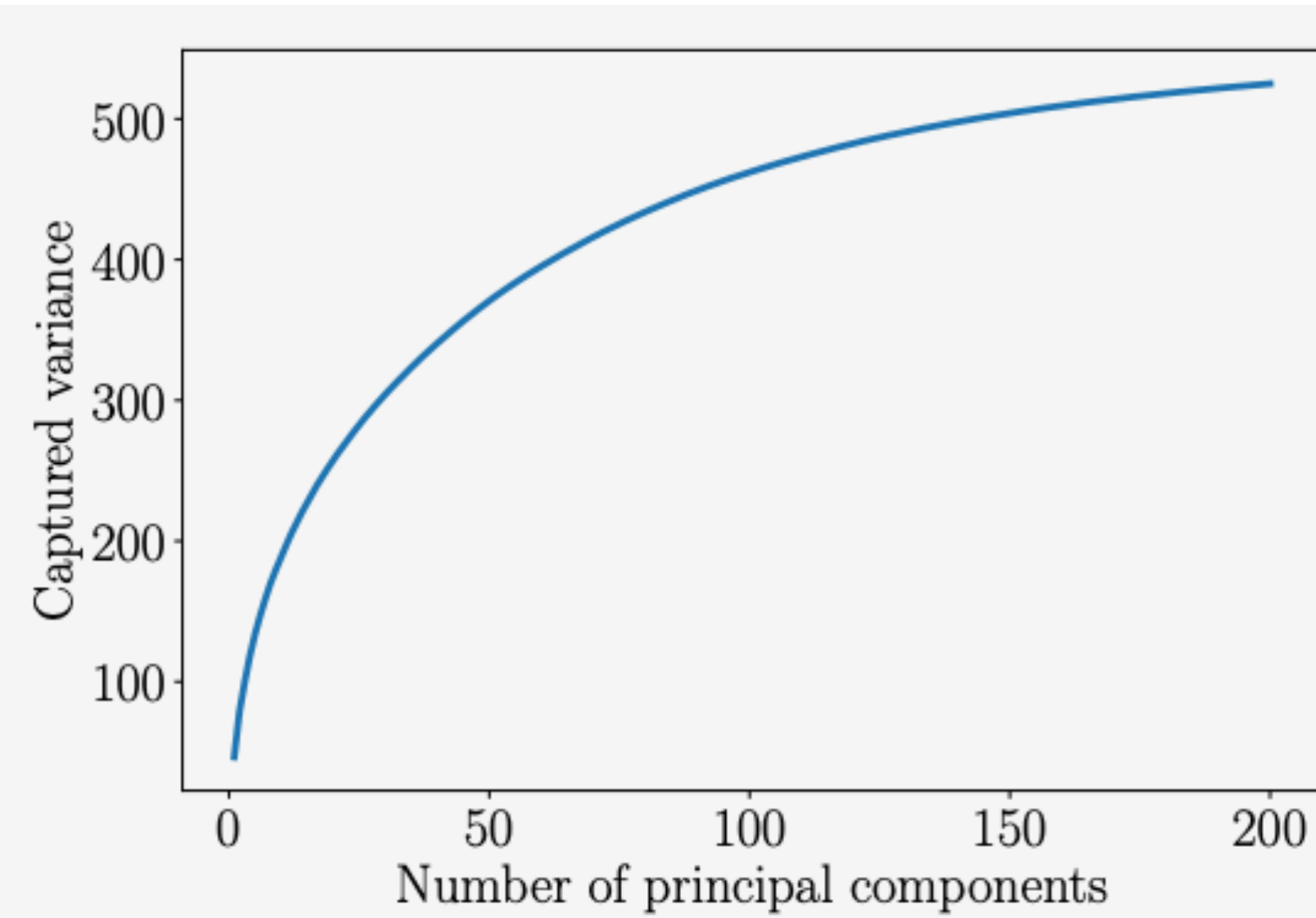
- 60,000 examples of handwritten digits 0 through 9.
- Each digit is a grayscale image of size 28×28, i.e., it contains 784 pixels.
- We can interpret every image in this dataset as a vector  $x \in \mathbb{R}^{784}$



# Example - PCA captured variance



(a) Top 200 largest eigenvalues



(b) Variance captured by the principal components.

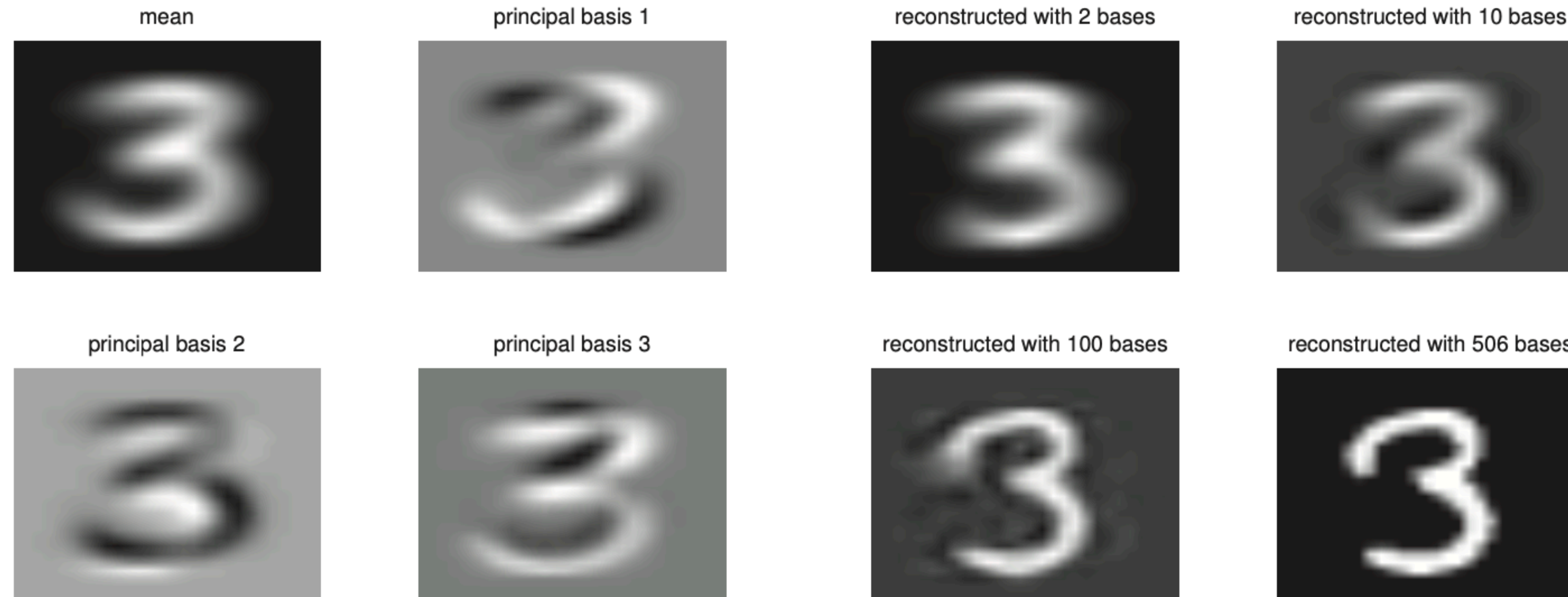
A 784-dim vector is used to represent an image

Taking all images of “3” in MNIST, we compute the eigenvalues of the data covariance matrix.

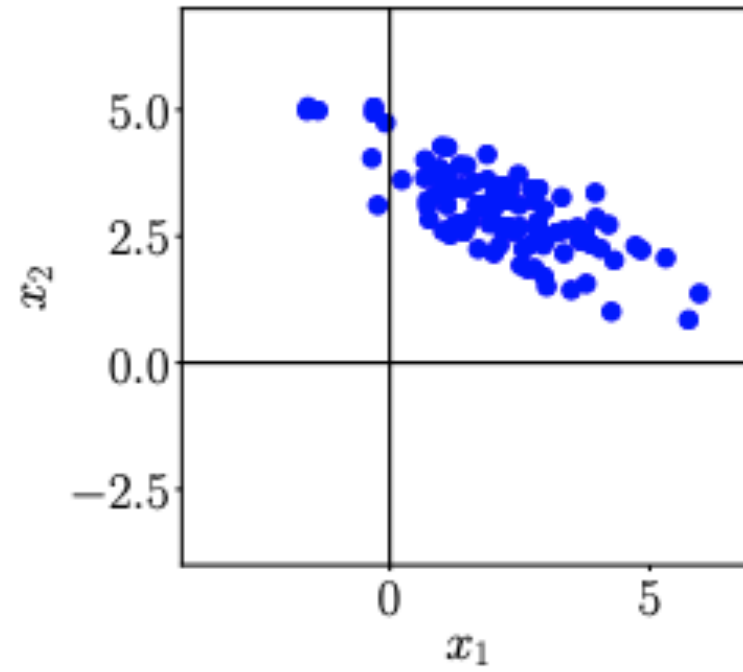
We see that only a few of them have a value that differs significantly from 0.

Most of the variance, when projecting data onto the subspace spanned by the corresponding eigenvectors, is captured by only a few principal components

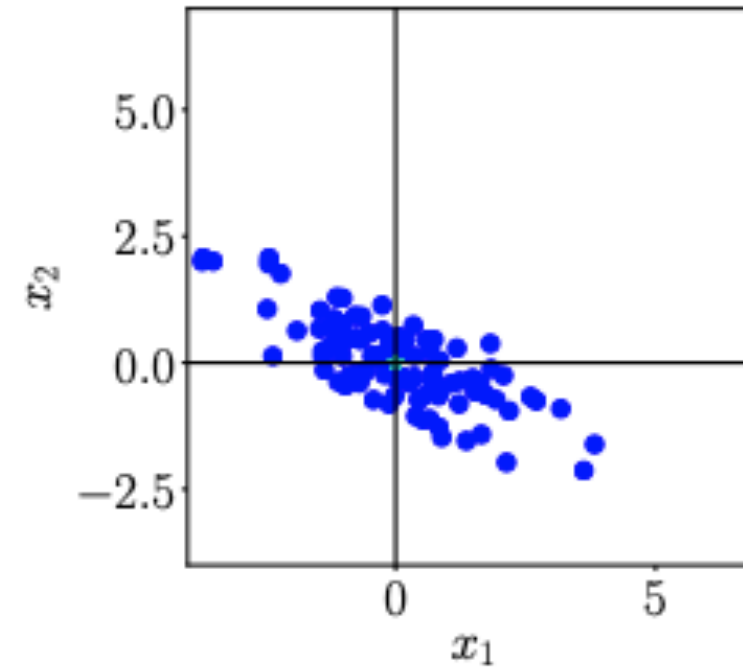
# Example - PCA reconstruction



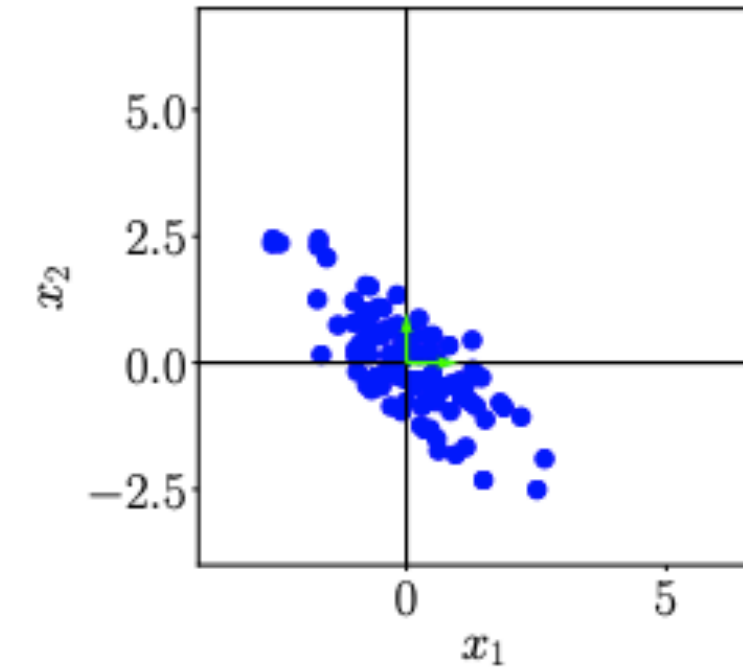
# PCA in practice



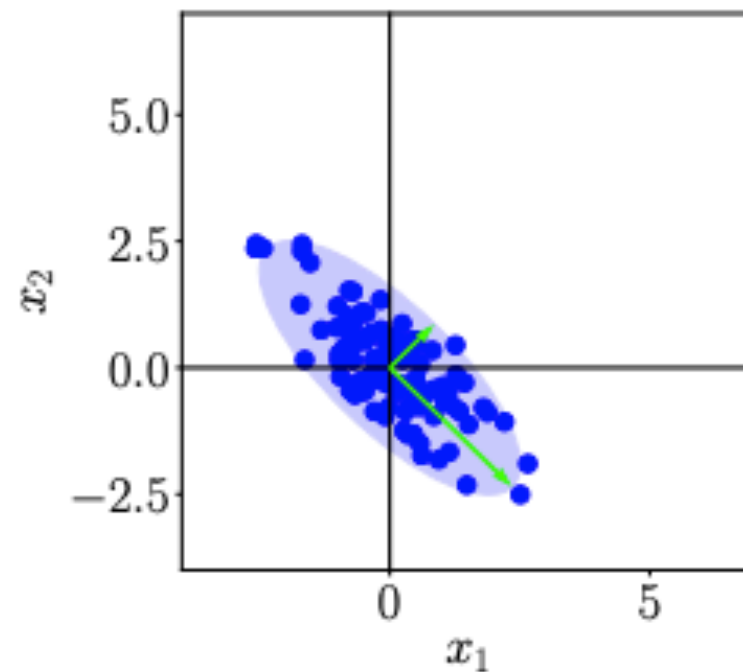
(a) Original dataset.



(b) Step 1: Centering by subtracting the mean from each data point.

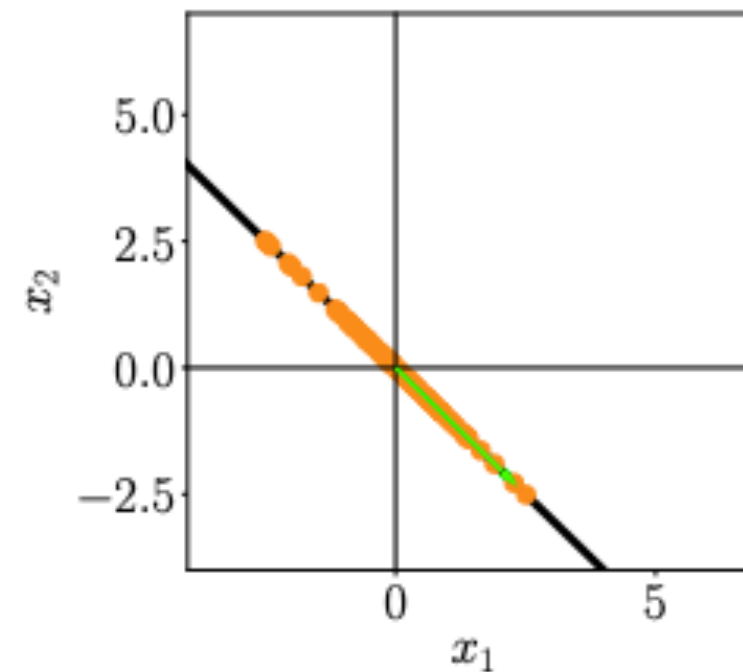


(c) Step 2: Dividing by the standard deviation to make the data unit free. Data has variance 1 along each axis.

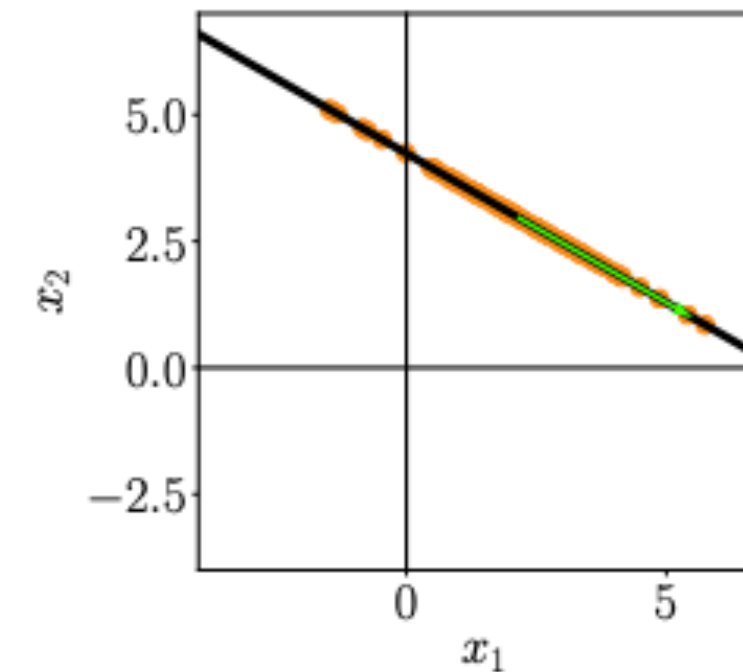


(d) Step 3: Compute eigenvalues and eigenvectors (arrows) of the data covariance matrix (ellipse).

eigendecomposition



(e) Step 4: Project data onto the principal subspace.



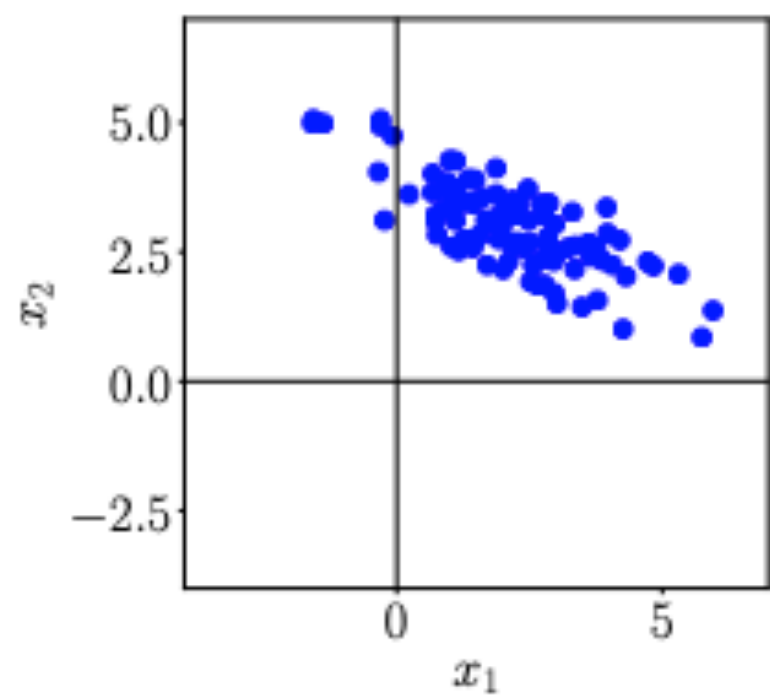
(f) Undo the standardization and move projected data back into the original data space from (a).

## Step 1. Mean subtraction

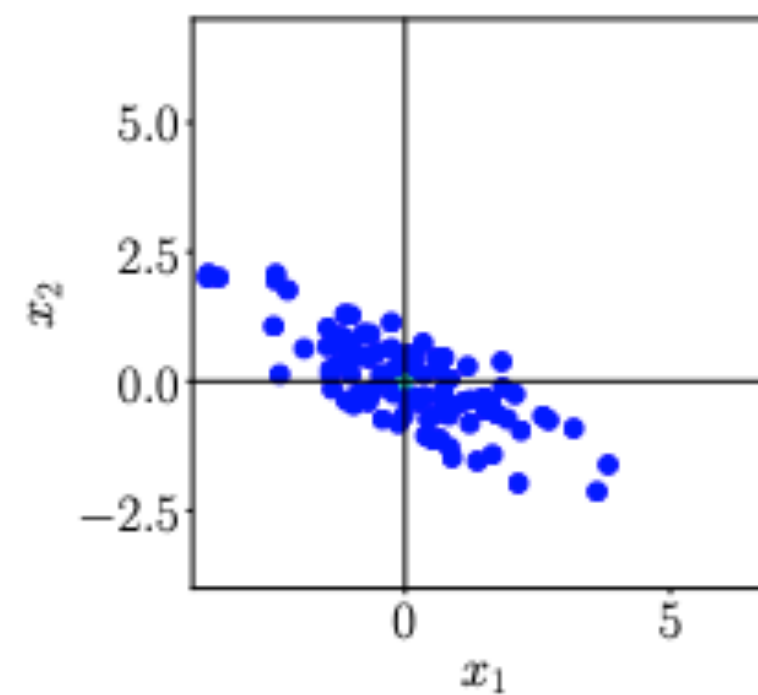
We center the data by computing the mean  $\mu$  of the dataset and subtracting it from every single data point. This ensures that the dataset has mean 0.

## Step 2. Standardisation

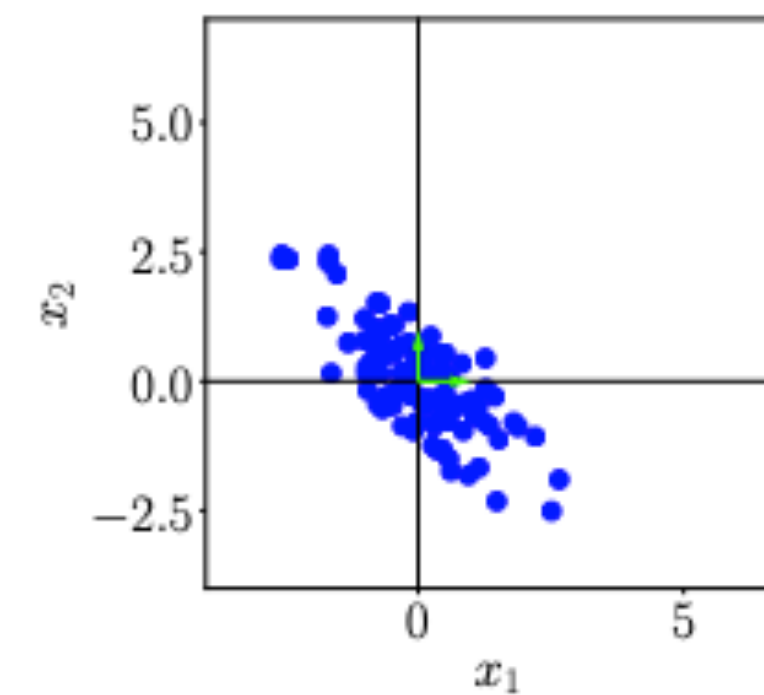
Divide the data points by the standard deviation  $\sigma_d$  of the dataset for every dimension. Now the data has variance 1 along each axis.



(a) Original dataset.



(b) Step 1: Centering by subtracting the mean from each data point.

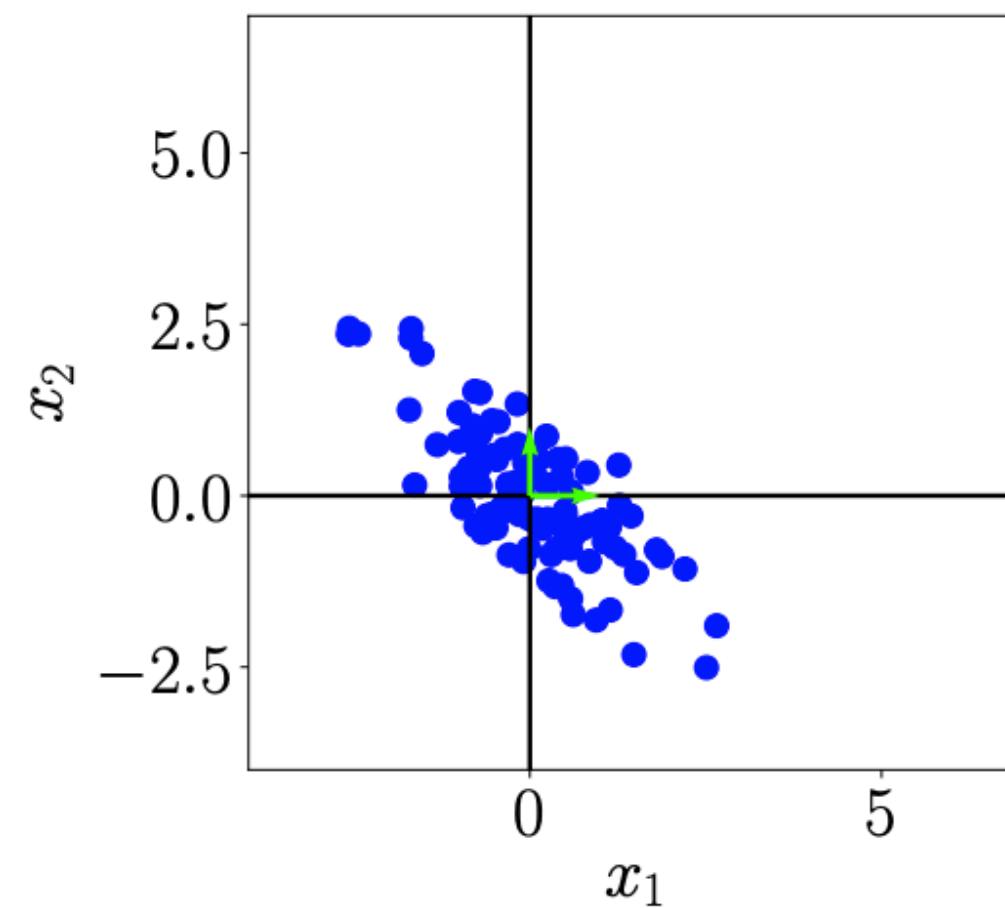


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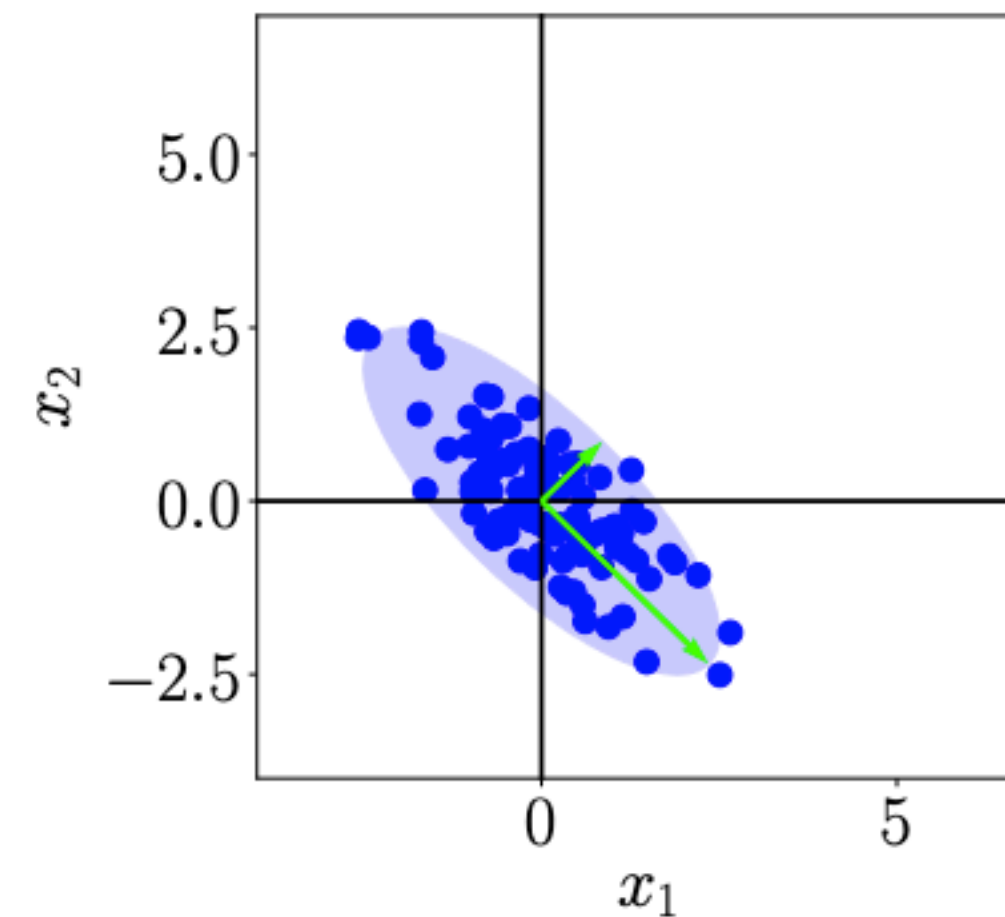


### Step 3. Eigendecomposition of the covariance matrix

Compute the data covariance matrix and its eigenvalues and corresponding eigenvectors. The longer vector (larger eigenvalue) spans the principal subspace  $U$



(c) Step 2: Dividing by the standard deviation to make the data unit free. Data has variance 1 along each axis.



(d) Step 3: Compute eigenvalues and eigenvectors (arrows) of the data covariance matrix (ellipse).

## 4. Projection

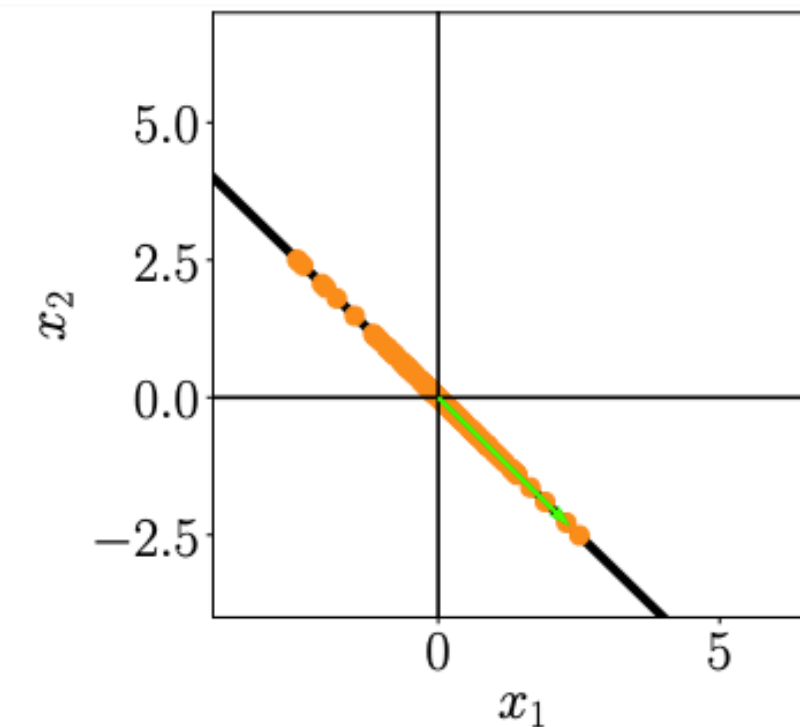
We can project any data point  $\mathbf{x}_* \in \mathbb{R}^D$  onto the principal subspace.

projection as  $\tilde{\mathbf{x}}_* = \mathbf{B}\mathbf{B}^T\mathbf{x}_*$

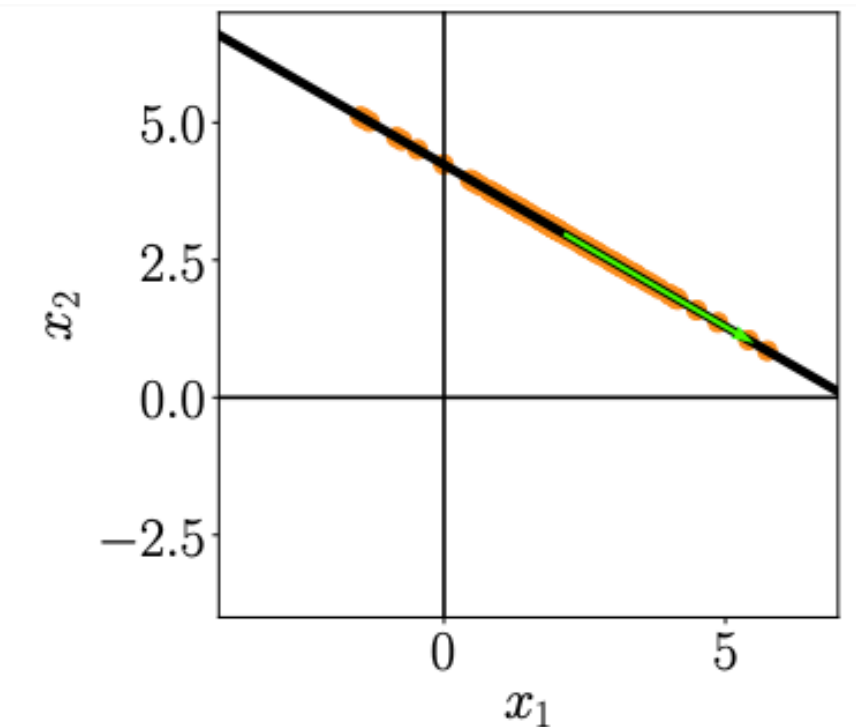
coordinates  $\mathbf{z}_* = \mathbf{B}^T\mathbf{x}_*$  with respect to the basis of the principal subspace. Here,  $\mathbf{B}$  is the matrix that contains the eigenvectors that are associated with the largest eigenvalues of the data covariance matrix as columns.

## 5. Rescaling data

To obtain our projection in the original data space (i.e., before standardization), we need to undo the standardization: multiply by the standard deviation before adding the mean.



(e) Step 4: Project data onto the principal subspace.



(f) Undo the standardization and move projected data back into the original data space from (a).

Exercise: Show that PCA is rotationally invariant

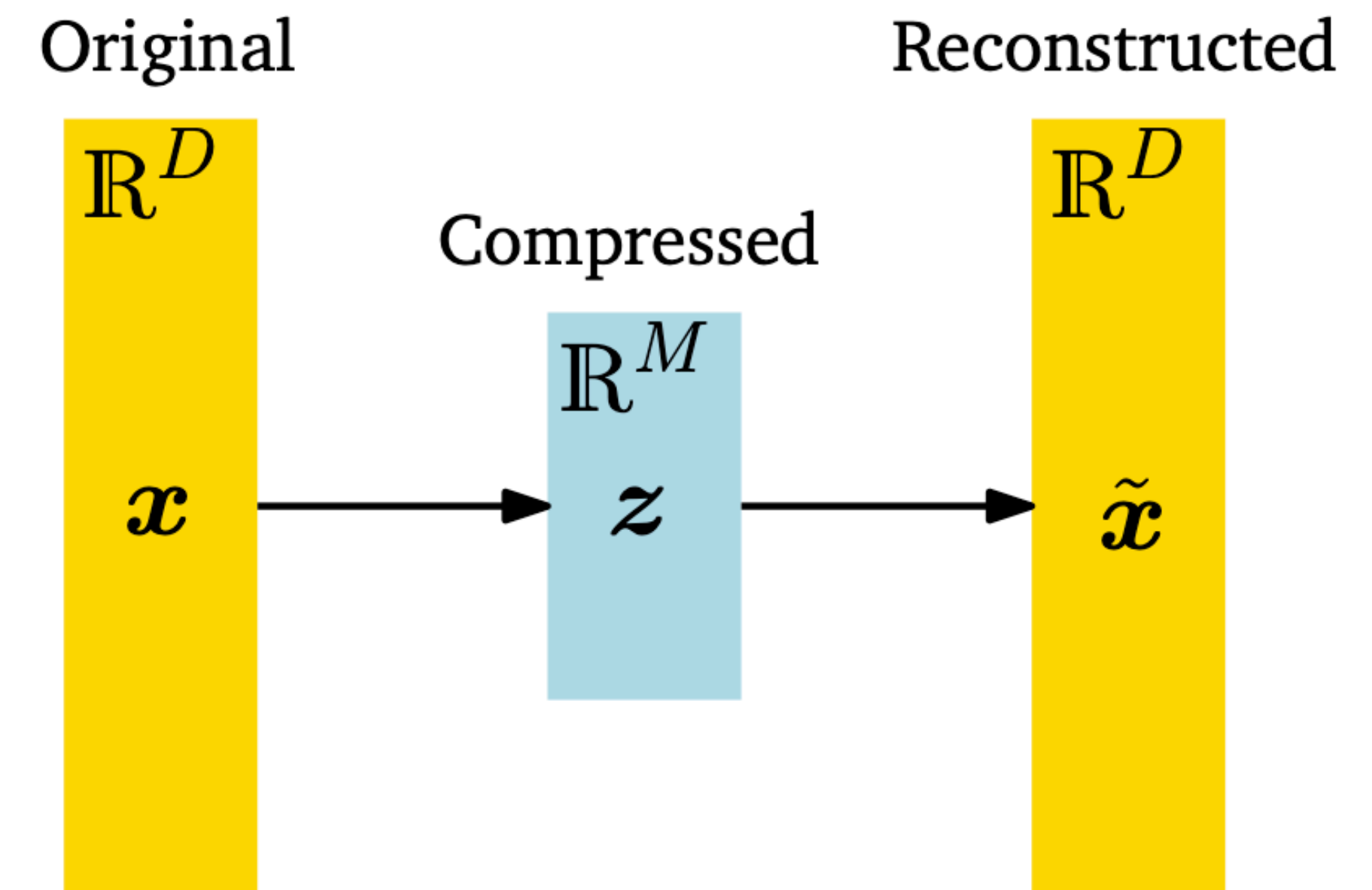
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# Problem setup



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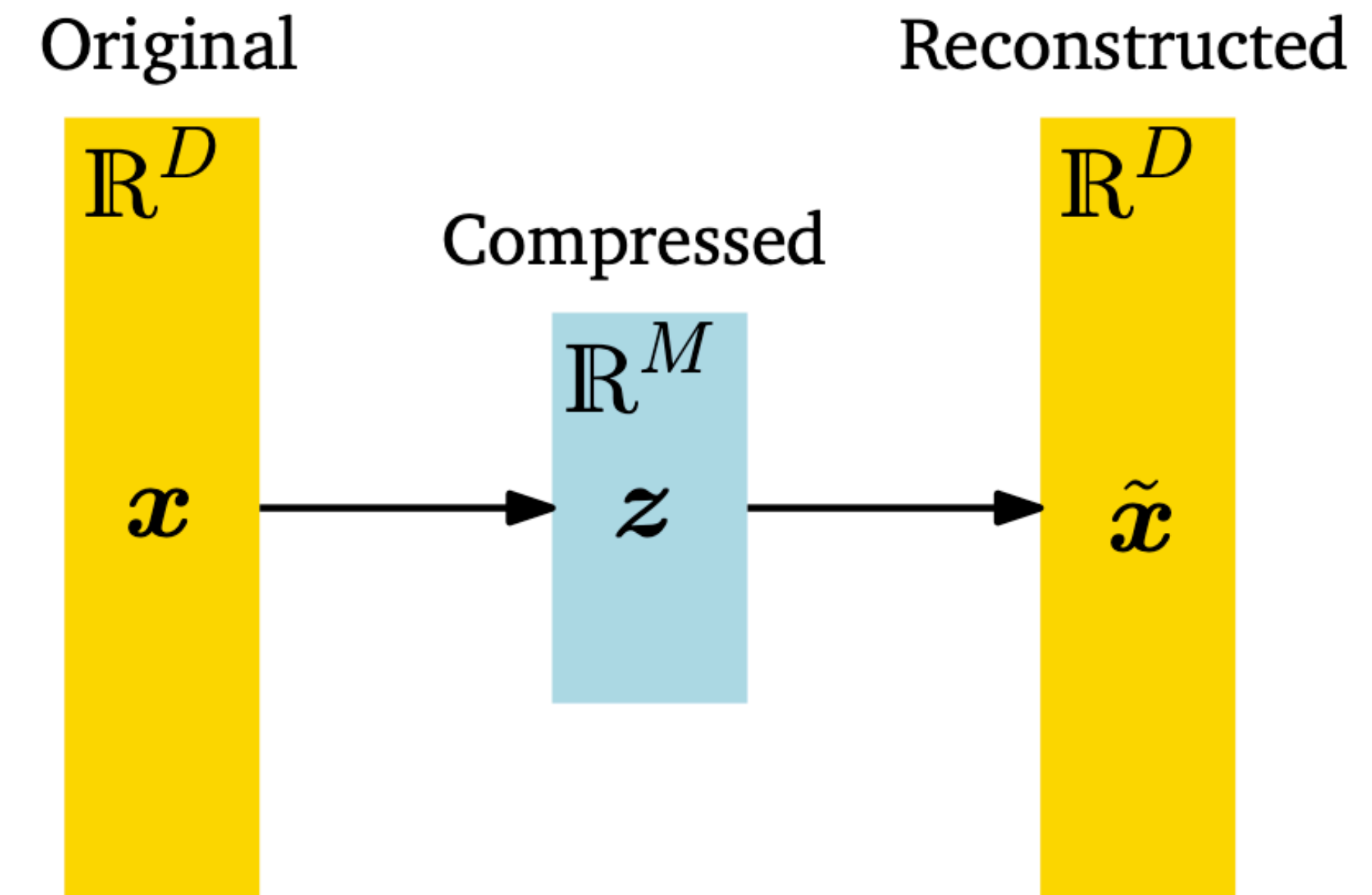
We consider an i.i.d. dataset  $X = \{x_1, x_2, \dots, x_N\}$ ,  $x_n \in \mathbb{R}^D$ ,

with mean  $\mathbf{0}$  and covariance matrix  $S = \frac{1}{N} \sum_{n=1}^N x_n x_n^\top$

We assume there exists a *low-dimensional* compressed representation (code):  $z_n = B^\top x_n$ ,  $z_n \in \mathbb{R}^M$ ,  $M < D$ .

The projection matrix:  $B = [b_1, b_2, \dots, b_M] \in \mathbb{R}^{D \times M}$ , columns are orthonormal.

*Reconstruction* using  $B$ :  $\tilde{x}_n = B z_n$



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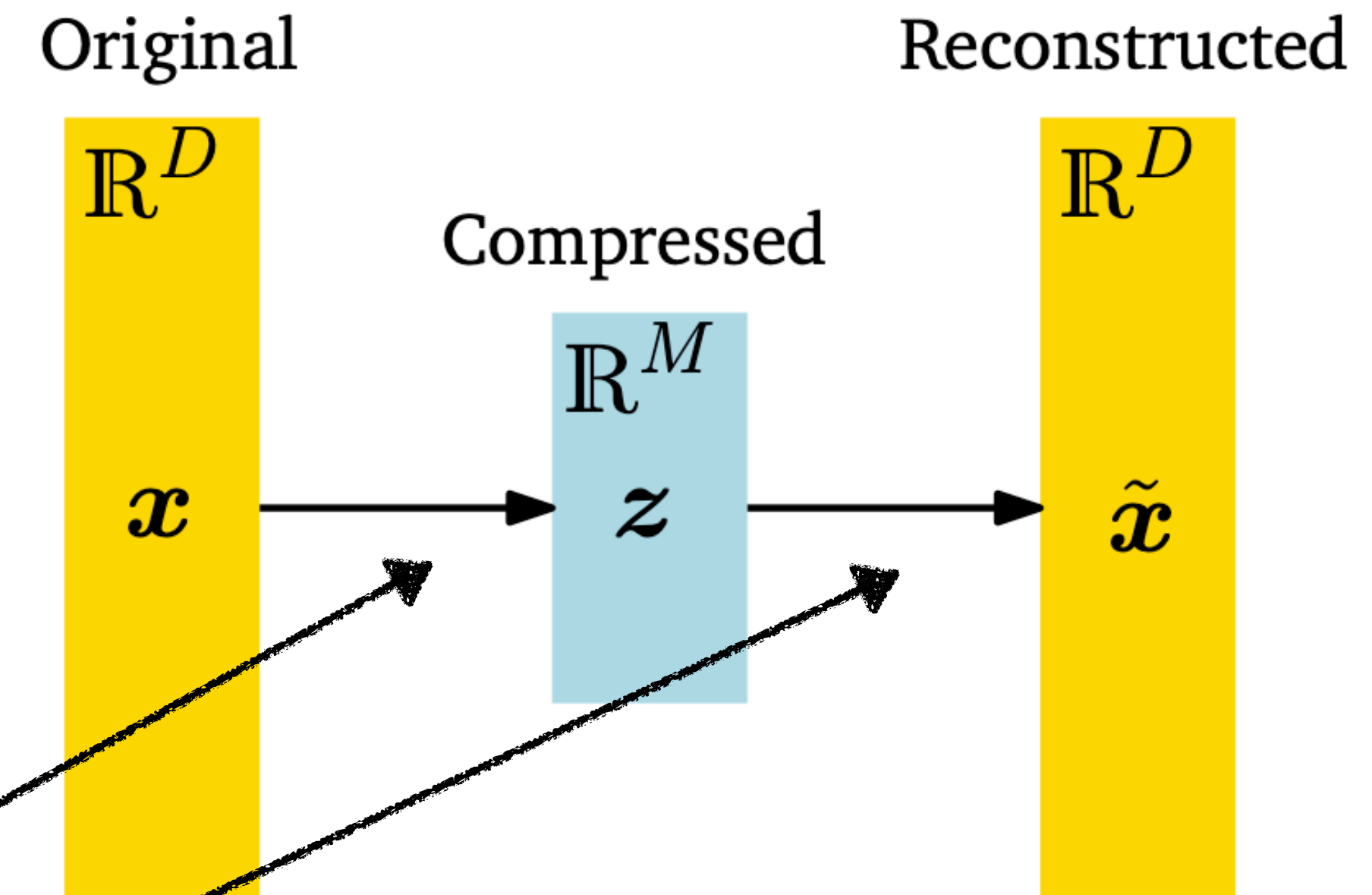
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**PCA: linear mappings**

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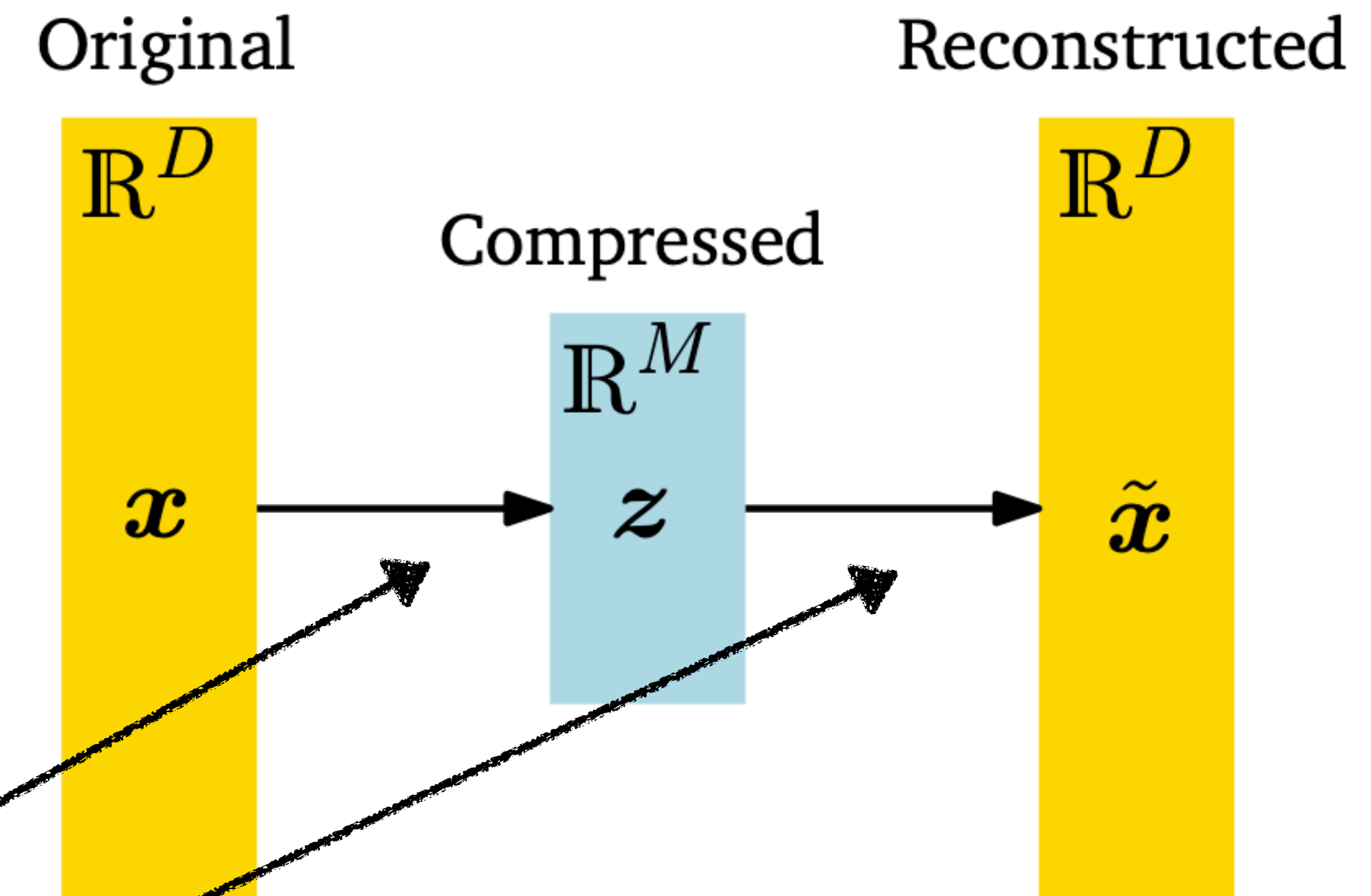
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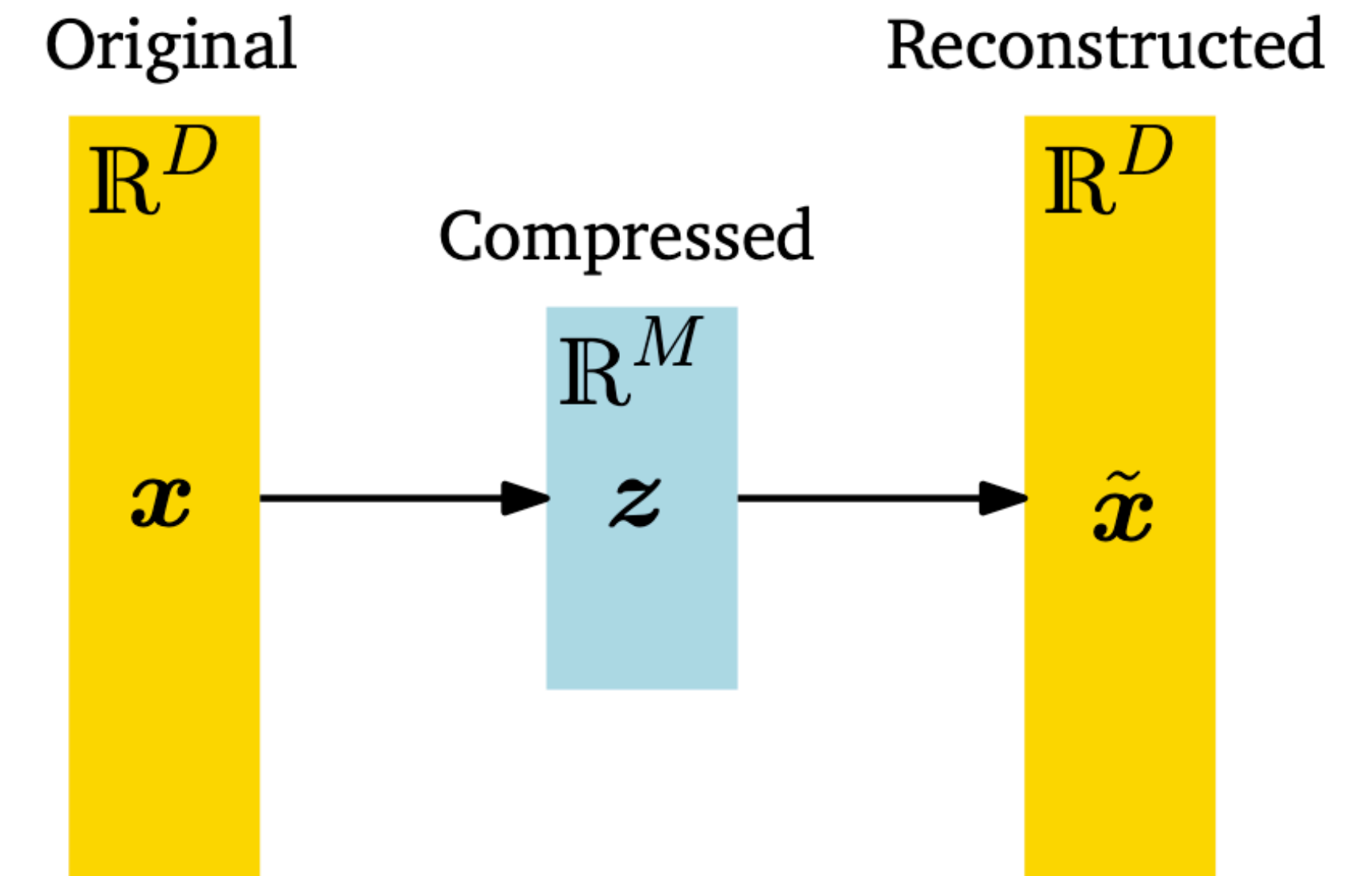
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**PCA: linear mappings**

**Goal:** find  $z_n$  and the *basis vectors*  $b_1, b_2, \dots, b_M$  so that the reconstructed data are *similar* to the original data, and the compressed data retain most of the *variation* in the original data

# PCA - two perspectives



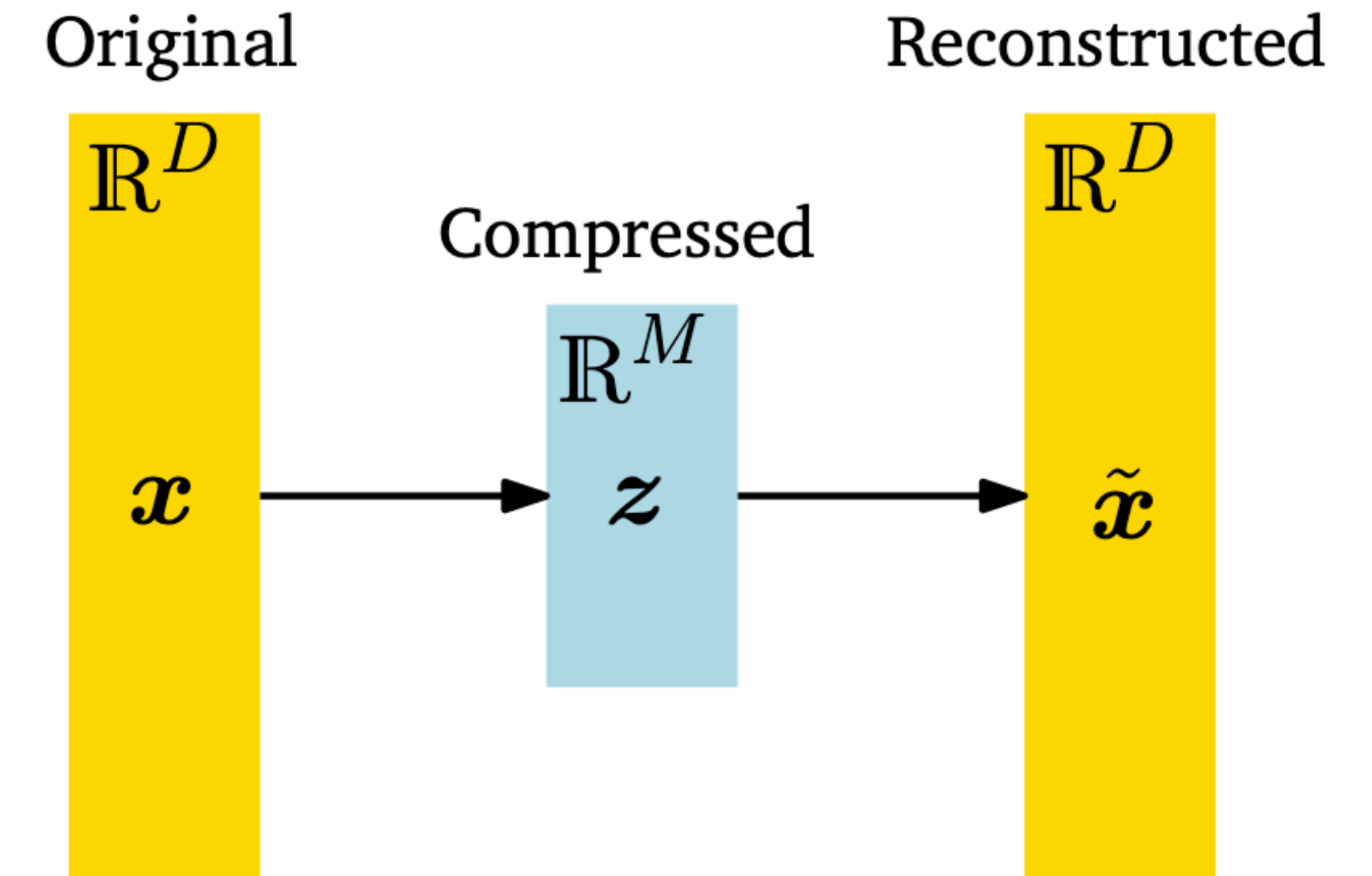
**PCA: linear mappings**

$$z_n = B^\top x_n, z_n \in \mathbb{R}^M, M < D$$

$$\tilde{x}_n = B z_n$$

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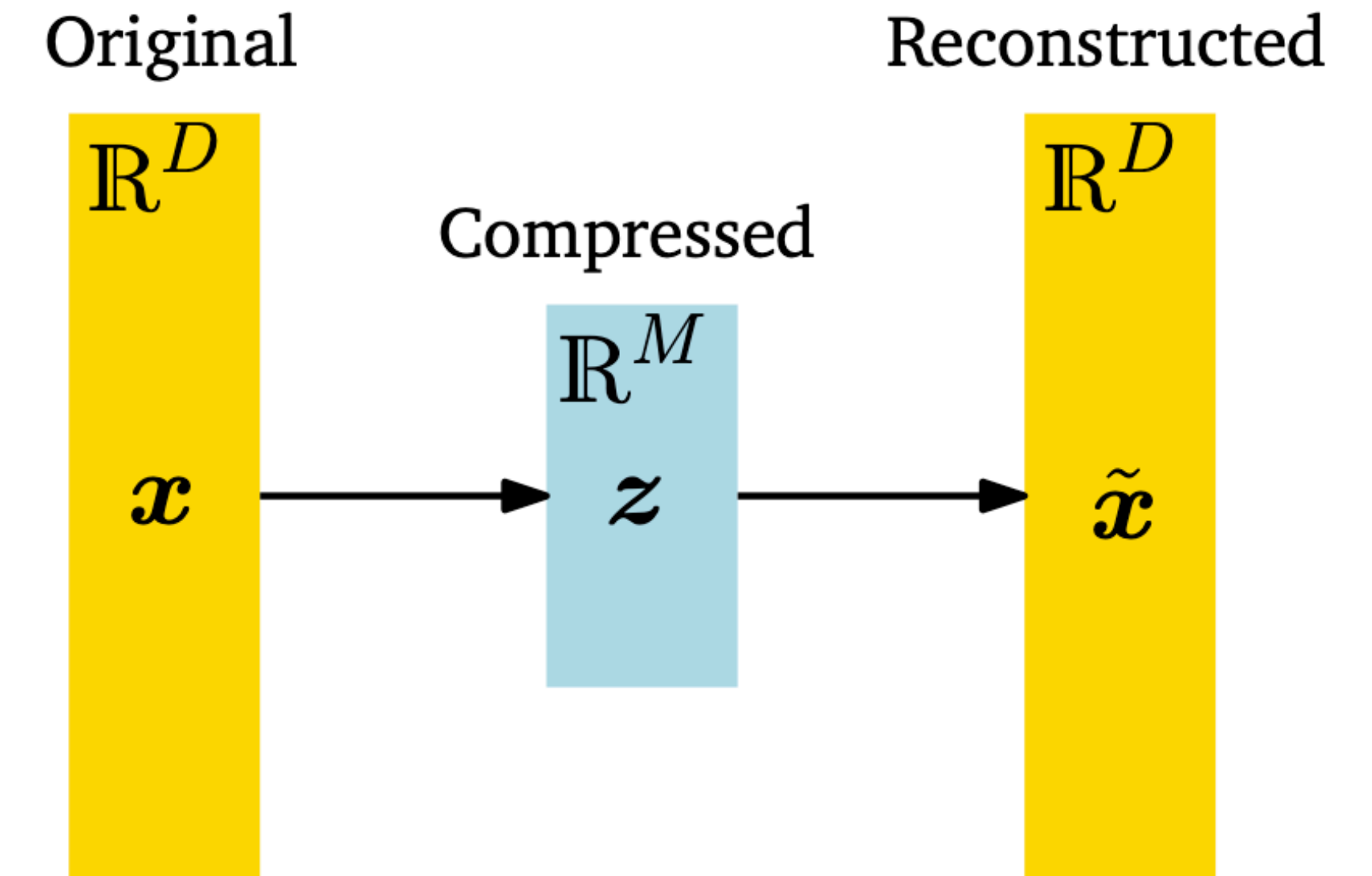
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**Question:** Next steps? Ideas?



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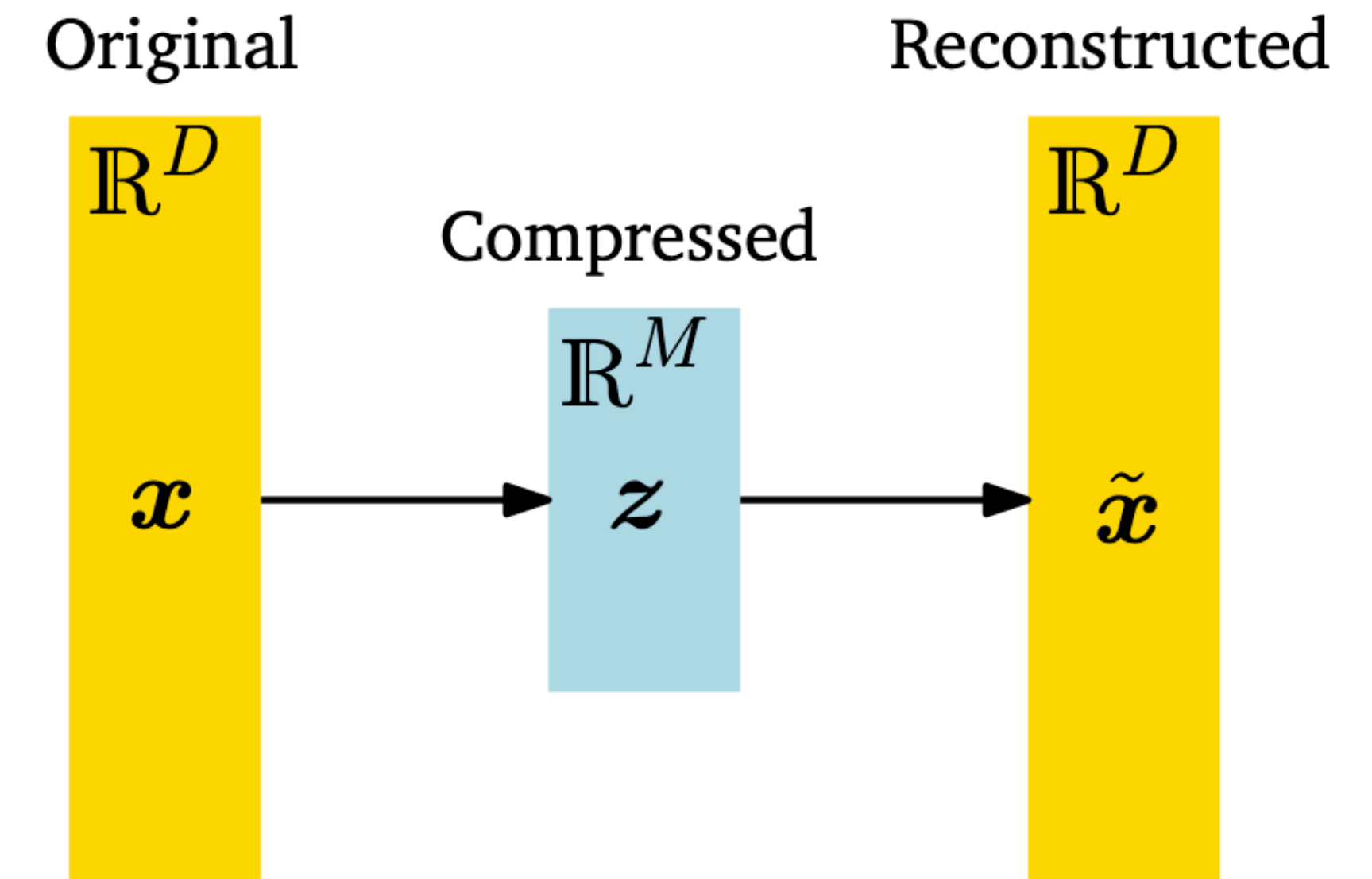
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**Question:** Next steps? Ideas?

**Answer:** Two approaches

- + Search for  $B$  that **maximises the variance** of the low-dimensional representations [analysis/max var perspective]
- + Search for  $B$  and  $z$  that minimises the reconstruction loss [synthesis/projection perspective]

Both give *identical* solutions! **Why?**



**PCA: linear mappings**

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2. PCA review
- 3. Linear, Gaussian latent variable models and GPLVM [whiteboard]**

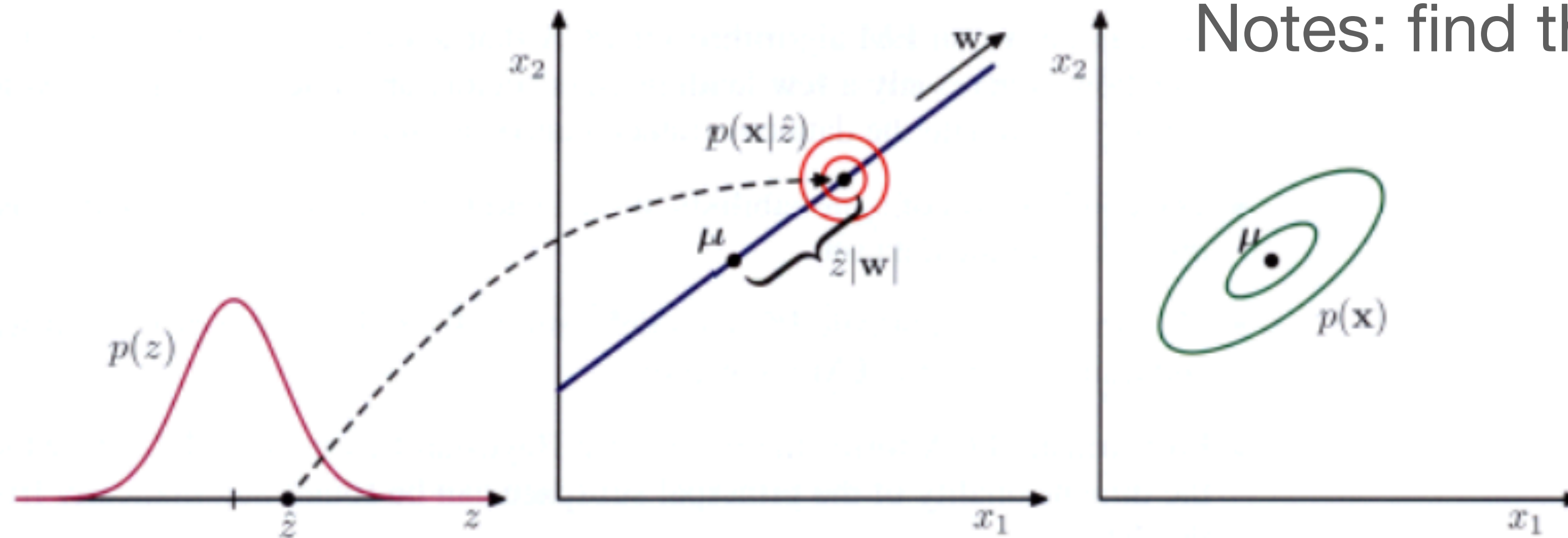
Reading: Bishop 12.1, 12.2, 12.4.2

# Probabilistic PCA

$$p(\mathbf{z}) = \mathcal{N}(\mathbf{z}; 0, \mathbf{I})$$
$$p(\mathbf{x} | \mathbf{z}) = \mathcal{N}(\mathbf{x}; W\mathbf{z} + \mu, \sigma^2 \mathbf{I})$$

- Benefits:
- + a proper probabilistic density model
  - + EM algorithm for computational efficiency
  - + Can be extended to handle binary/categorical data
  - + Can be extended to handle discrete latent variables
  - + Generate samples

Notes: find the same principal subspace as PCA



**Figure 12.9** An illustration of the generative view of the probabilistic PCA model for a two-dimensional data space and a one-dimensional latent space. An observed data point  $\mathbf{x}$  is generated by first drawing a value  $\hat{z}$  for the latent variable from its prior distribution  $p(z)$  and then drawing a value for  $\mathbf{x}$  from an isotropic Gaussian distribution (illustrated by the red circles) having mean  $w\hat{z} + \mu$  and covariance  $\sigma^2 \mathbf{I}$ . The green ellipses show the density contours for the marginal distribution  $p(\mathbf{x})$ .

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3. Probabilistic PCA: linear, Gaussian latent variable models and GPLVM

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Enjoy the break and see you in W12!