

# Unsupervised learning - introduction : Clustering

## Unsupervised learning

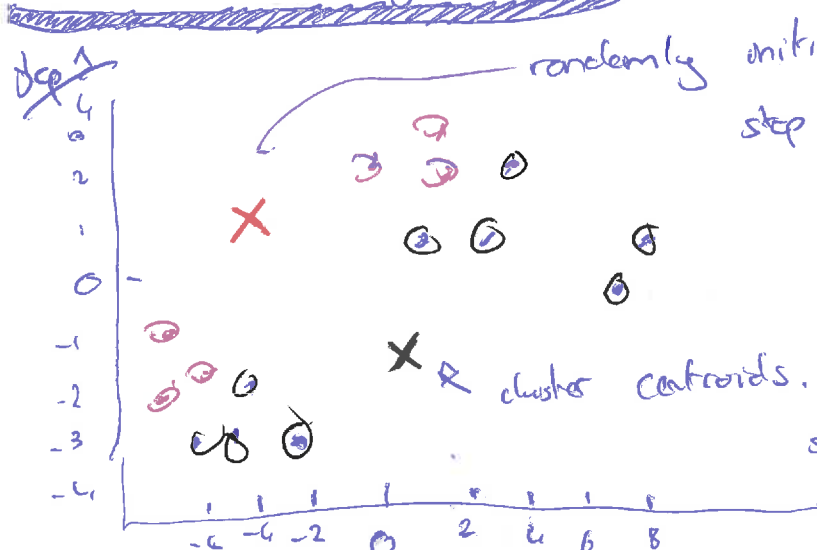


Training set:  $\{x^{(1)}, x^{(2)}, x^{(3)}, \dots, x^{(n)}\}$

## Applications of clustering

- Market segmentation
- Social network Analysis
- Organise computing clusters
- astronomical Analysis

## K-means Algorithm



randomly initialize  $k$  cluster center points  
step 1) Cluster assignment step

so if a data point is closer to a cluster centroid  $\rightarrow$  it is assigned to that cluster centroid.

step 2) move centroid step

compute average of all the assigned data points of a cluster and move the centroid to this new centrepaint.



now closer together - centroids "moved"

$\rightarrow$  after awhile the centroid locations no longer change.

# K-means algorithm

Input:

-  $k$  (number of clusters) ✓

- training set  $\{x^{(1)}, x^{(2)}, \dots\}$  ✓ (no  $y$ ...)

$x^{(i)} \in \mathbb{R}^n$  (drop  $x_0 = 1$ )

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## K-means algorithm

Randomly initialize  $k$  cluster centroids  $\mu_1, \mu_2, \dots, \mu_k \in \mathbb{R}^n$

Repeat { for  $i = 1$  to  $m$

cluster assignment step  $c^{(i)} := \text{index (from 1 to } k) \text{ of cluster centroid closest to } x^{(i)}$   
 $\min_k \|x^{(i)} - \mu_k\|_2 \text{ length.}$

for  $k = 1$  to  $k$

move centroid step.  $\mu_k := \text{average (mean) of points assigned to cluster } k$

↳ let assume this cluster (1) has the values  $x^{(1)}, x^{(5)}, x^{(6)}, x^{(10)}$  assigned to it.

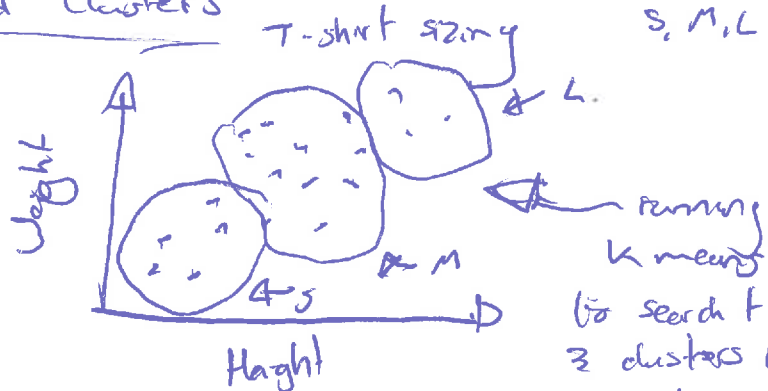
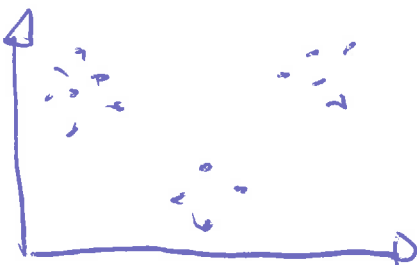
$\therefore c^{(1)} = 2, c^{(5)} = 2, c^{(6)} = 2, c^{(10)} = 2.$

cluster assignment step  
 distance between  $i$ th training example and cluster centroid  
 ↳ so, picking the cluster centroid  $c^{(i)}$  that is closest to our training example  $x^{(i)}$

$$\mu_2 = \frac{1}{4} [x^{(1)} + x^{(5)} + x^{(6)} + x^{(10)}] \in \mathbb{R}^n$$

notes: delete a cluster if it has no points assigned to it  
 ↳ or reinitialize it through randomization

## K-means for non-separated clusters



running K means  
 to search for 3 clusters for S, M, L.

# Clustering - optimization objective

## K-means optimization objective

$c^{(i)}$  = index of cluster  $(1, 2, \dots, K)$  to which example  $x^{(i)}$  is currently assigned.

$\mu_k$  = cluster centroid  $k \in \{1, 2, \dots, K\}$  ( $\mu_k \in \mathbb{R}^n$ )

$\mu_{c^{(i)}}$  = cluster centroid of cluster to which  $x^{(i)}$  has been assigned

$x^{(i)} \rightarrow 5 \therefore c^{(i)} = 5$

$x$  has been assigned to cluster 5

lower case  $k$  is an index into  $K = \{1, 2, \dots, K\}$

$\mu_{c^{(i)}} = \mu_5 = ?$

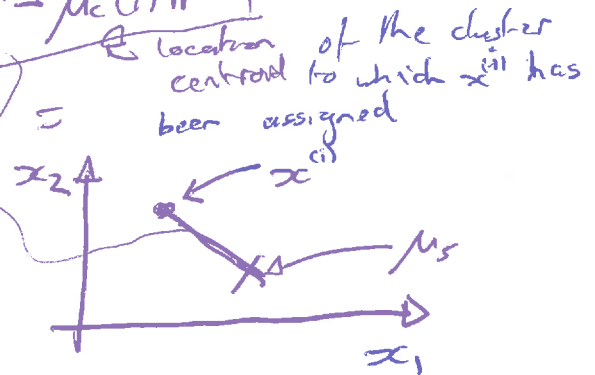
optimization objective:

$$J(c^{(1)}, \dots, c^{(m)}, \mu_1, \dots, \mu_K) = \frac{1}{m} \sum_{i=1}^m \|x^{(i)} - \mu_{c^{(i)}}\|^2$$

$$\min_{c^{(1)}, \dots, c^{(m)}, \mu_1, \dots, \mu_K} J(c^{(1)}, \dots, c^{(m)}, \mu_1, \dots, \mu_K)$$

$\mu_1, \dots, \mu_K$

→ Sometimes called the "distortion cost function"



→ The cluster assignment step in minimizing  $J(\dots)$  with respect to  $c^{(1)}, c^{(2)}$  etc. —  $c^{(m)}$  (holding  $\mu_1, \dots, \mu_K$  fixed)

→ minimizes  $J(\dots)$  w.r.t.  $\mu_1, \dots, \mu_K$ .

# Clustering - Random initialization

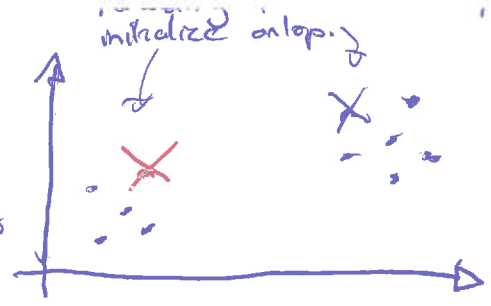
should have  $K < m$

Randomly pick  $K$  training examples

Set  $\mu_1, \dots, \mu_K$  equal to these  $K$  examples

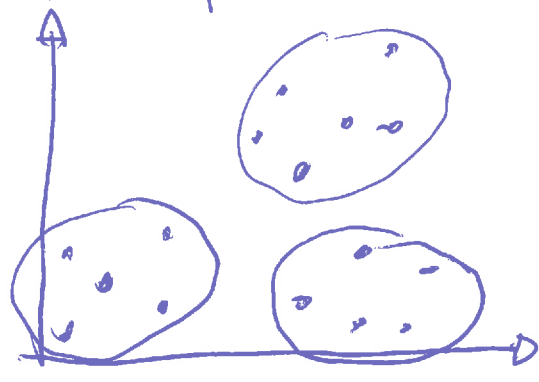
$$\begin{aligned} \mu_1 &= x^{(i)} \\ \mu_2 &= x^{(j)} \\ &\vdots \end{aligned}$$

lets say:  
 $K=2$   
two examples



$K$  means can converge to different solutions depending how the clusters were initialized

Local optima.  $\rightarrow$  can get stuck at different local optima.



initialize  $K$  means lds of times (randomly) to increase the chances of finding the global minimum.

## Random initialization

typical 50  $\rightarrow$  1000

for  $i = 1$  to 100 {

Randomly initialize  $K$ -means

Run  $K$  means to get  $c^{(1)}, \dots, c^{(m)}$

Compute cost function (distortion)  $\mu_1, \dots, \mu_K$

$$J(c^{(1)}, \dots, c^{(m)}, \mu_1, \dots, \mu_K)$$

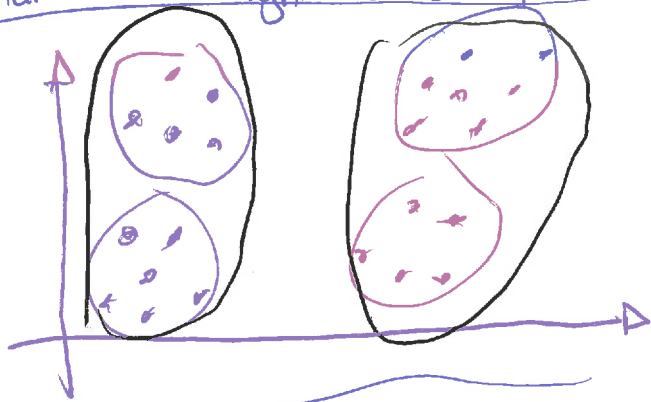
Pick clustering that gives lowest cost  $J(c^{(1)}, \dots, c^{(m)}, \mu_1, \dots, \mu_K)$

$K=2 \rightarrow 10$

If you have  $K > 10$ , multiple random initializations will not make much difference

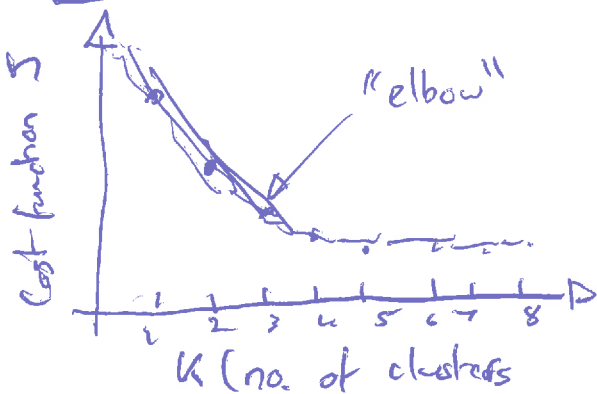
# clustering - choosing the number of clusters

What is the right value of  $K$ ?

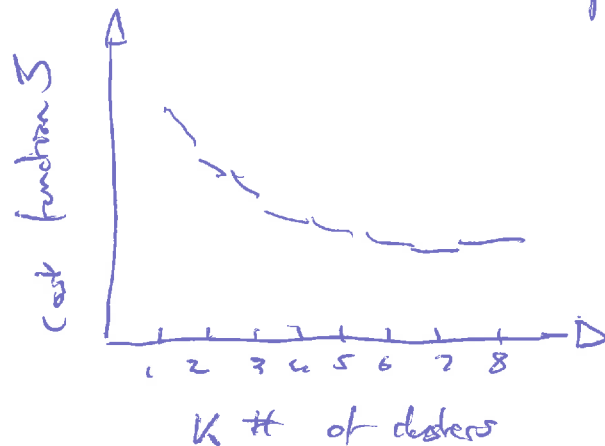


choosing the value of  $K$

Elbow method: "worth a shot"



Sometimes this happens



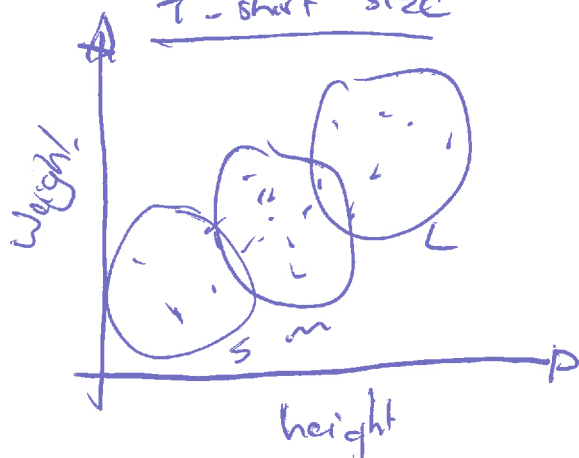
Choosing the value of  $K$

Sometimes, you're running  $K$  means to get clusters for some later/ downstream purpose. Evaluate  $K$ -means based on a metric for how well it performs for that later purpose.

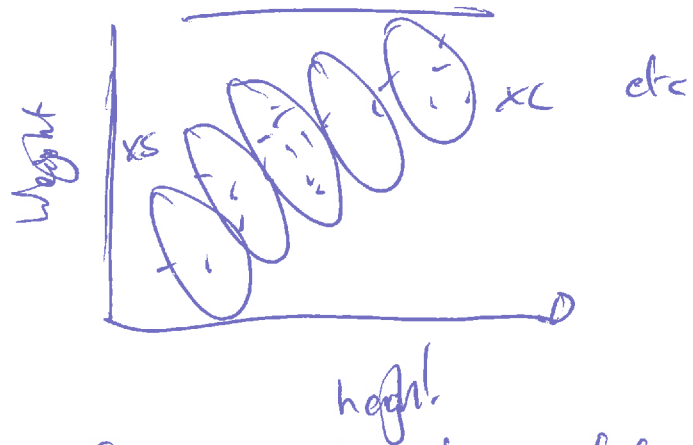
$K=3$

S, M, L

T-shirt size



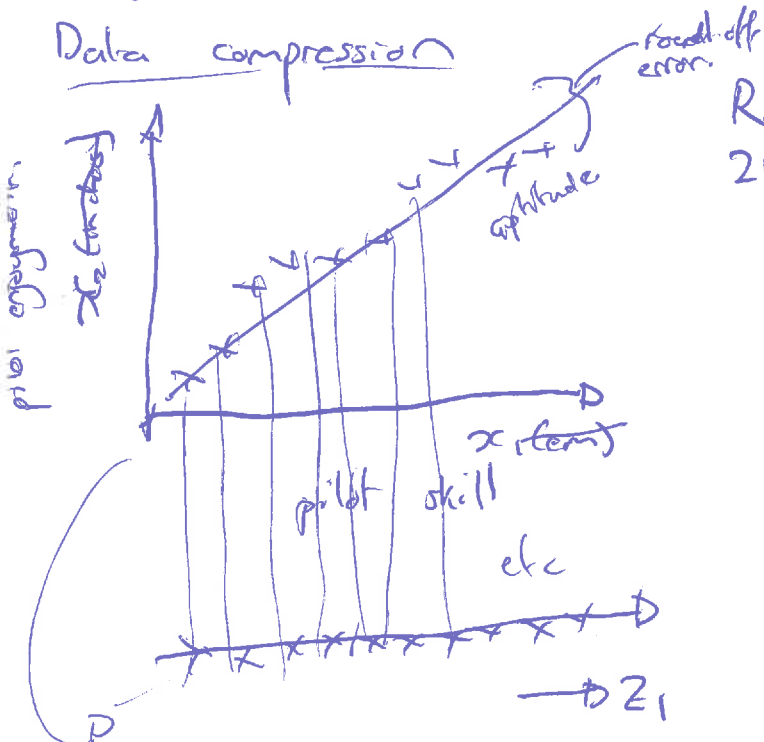
$K=5$  T-shirt size XS, S, M, L, XL



deciding what set of sizes for t-shirts to manufacture is

# Dimensionality Reduction - Motivation I: data compression

## Data compression



Reduce data from 2D to 1D

$$x^{(1)} \in \mathbb{R}^2 \rightarrow z^{(1)} \in \mathbb{R}$$

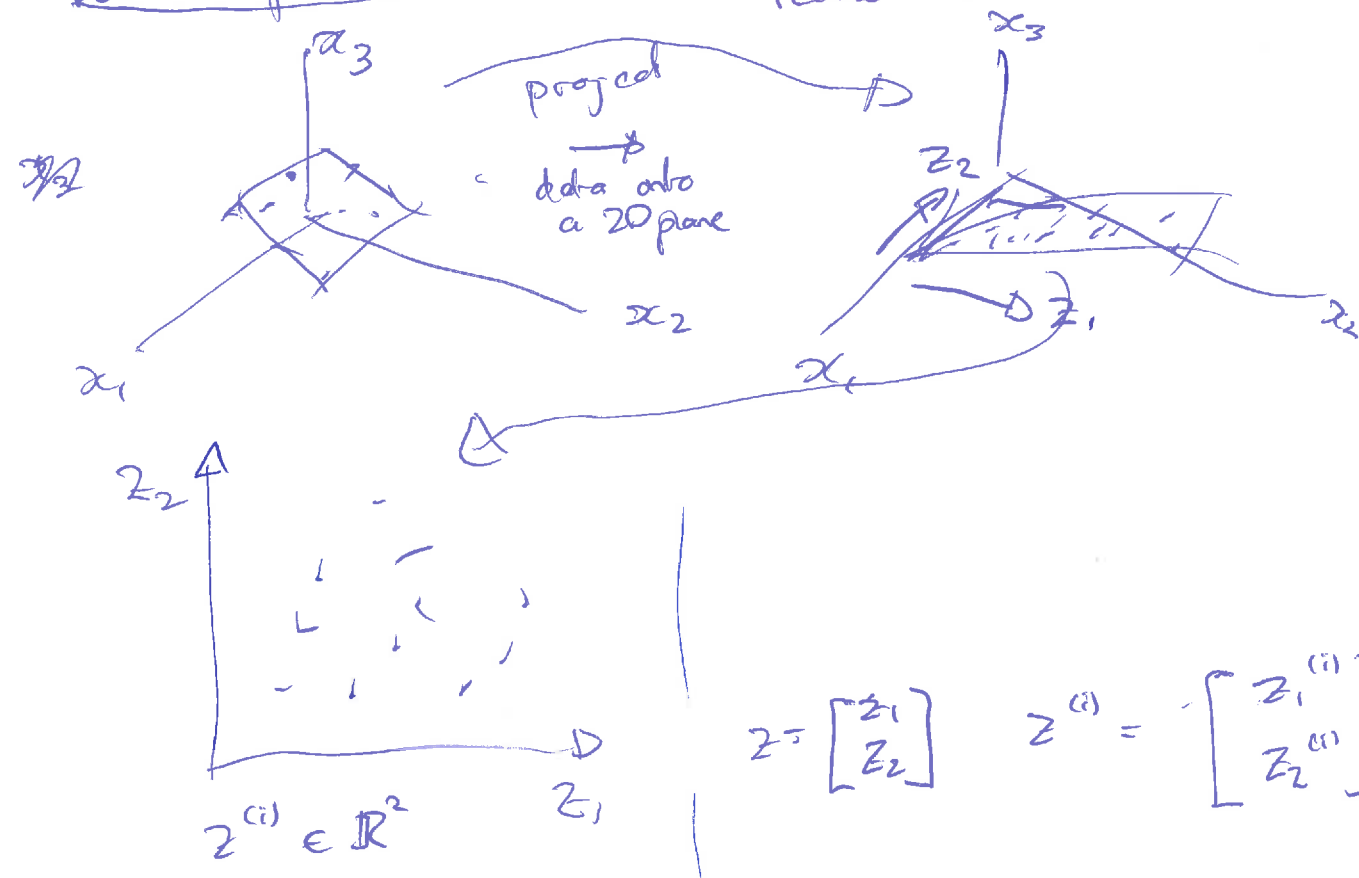
$$x^{(2)} \in \mathbb{R}^2 \rightarrow z^{(2)} \in \mathbb{R}$$

$$\vdots$$

$$x^{(n)} \rightarrow z^{(n)}$$

## Data Compression

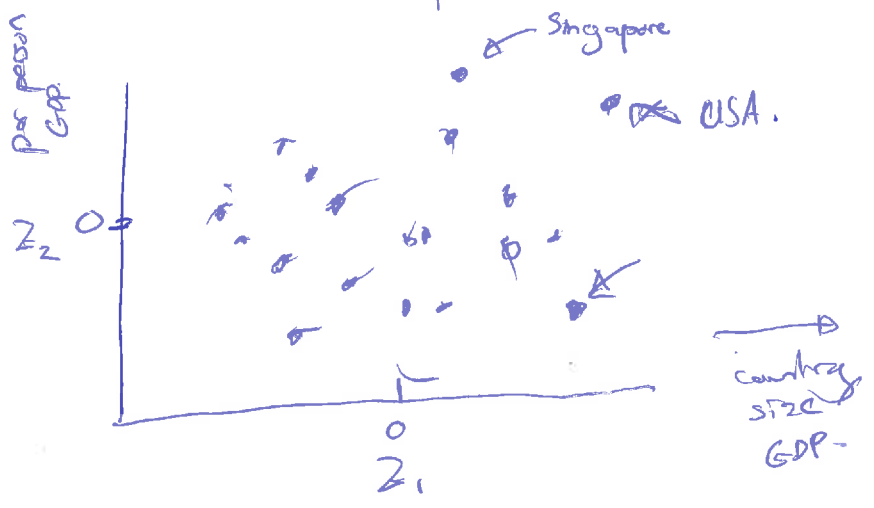
1000D to 100D  
reduce data from 3D to 2D.



$$Z = \begin{bmatrix} z_1 \\ z_2 \end{bmatrix} \quad Z^{(i)} = \begin{bmatrix} z_1^{(i)} \\ z_2^{(i)} \end{bmatrix}$$

# Dimensionality Reduction 3 Motivation II 2 data visualization

Reduce data from 80D to 2D.



# Dimensionality Reduction - principal component analysis algorithm

## Data preprocessing

Training set:  $x^{(1)}, x^{(2)}, \dots, x^{(m)}$

preprocessing (feature scaling / mean normalization)

$$\mu_j = \frac{1}{m} \sum_{i=1}^m x_j^{(i)}$$

Replace each  $x_j^{(i)}$  with  $x_j - \mu_j$

If different features on different scales (e.g.,  $x_1$  = size of house,  $x_2$  = number of bedrooms), scale features to have comparable range of values.

$$x_j^{(i)} \mapsto \frac{x_j^{(i)} - \mu_j}{s_j}$$

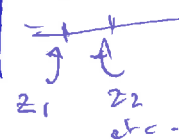
$s_j$   $\leftarrow$  standard deviation of feature  $j$ .

## Principal Component Analysis Algorithm

- Reduce data from 2D to 1D

- Reduce data from 3D to 2D.

$$x^{(i)} \in \mathbb{R}^2 \rightarrow z^{(i)} \in \mathbb{R}$$
$$Z = \begin{bmatrix} z_1 \\ z_2 \end{bmatrix}$$

e.g.  


## Principal Component Analysis (PCA) Algorithm

Reduce data from  $n$ -dimensions to  $k$ -dimensions

Compute "covariance matrix"

not to be confused with summation

$$\Sigma = \frac{1}{m} \sum_{i=1}^m (x^{(i)})(x^{(i)})^T$$

$n \times 1$     $1 \times n$   $\rightarrow$   $n \times n$  matrix

Compute "eigenvectors" of matrix  $\Sigma$   $\rightarrow$   $n \times n$  matrix

$$\rightarrow [U, S, V] = \text{svd}(\Sigma) \quad (\text{or eig}(\Sigma))$$

$\leftarrow$  octave    $\leftarrow$  singular value decomposition

$$U = \begin{bmatrix} | & | & | & \dots & | \\ u^{(1)} & u^{(2)} & u^{(3)} & \dots & u^{(n)} \\ | & | & | & \dots & | \end{bmatrix}$$

$\underbrace{\hspace{10em}}_K$

vectors for plane

$$U \in \mathbb{R}^{n \times n}$$

$$u^{(1)}, \dots, u^{(k)}$$

$k$  direction onto which we want to project the data.



# Dimensionality reduction = principal component analysis algorithm

from  $[U, S, V] = \text{svd}(\text{Sigma})$ , we get:

$$U = \begin{bmatrix} | & | & & | \\ u^{(1)} & u^{(2)} & \dots & u^{(n)} \\ | & | & & | \end{bmatrix} \in \mathbb{R}^{n \times n}.$$

K.

$$X \in \mathbb{R}^n \rightarrow Z \in \mathbb{R}^k$$

$$Z = \begin{bmatrix} | & | & & | \\ u^{(1)} & u^{(2)} & \dots & u^{(k)} \\ | & | & & | \end{bmatrix}^T$$

n x k.

$U_{\text{reduce}}$

$$X^{(i)} = \begin{bmatrix} u^{(1)} \\ u^{(2)} \\ \vdots \\ u^{(k)} \end{bmatrix} \times \begin{bmatrix} \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \end{bmatrix} \times \begin{bmatrix} \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \end{bmatrix}$$

k x n

$$Z \in \mathbb{R}^k$$

## PCA Algorithm summary

$$\text{Sigma} = \frac{1}{m} \sum_{i=1}^m (x^{(i)})(x^{(i)})^T \quad \text{implemented on octave looks like} \rightarrow \text{Sigma} = (1/m) * X' * X;$$

$$\left. \begin{aligned} [U, S, V] &= \text{svd}(\text{Sigma}); \\ U_{\text{reduce}} &= U(:, 1:k); \\ Z &= U_{\text{reduce}}' * X; \end{aligned} \right\} \text{ octave code}$$

$x \in \mathbb{R}^n$   ~~$x_0 = 1$~~

# Dimension Reduction : Reconstruction from compressed representation

in the previous examples we ~~app~~ reduced a 2D (or  $N^{\text{th}}$  D) dimension space, down to 1D using

$$(x_2, x_1) \left\{ \begin{array}{l} \underset{\substack{\uparrow \\ \text{new} \\ \text{dimension (1D)}}}{Z} = U^T \text{reduce } x \end{array} \right.$$

using to reverse it, we need to do the following:

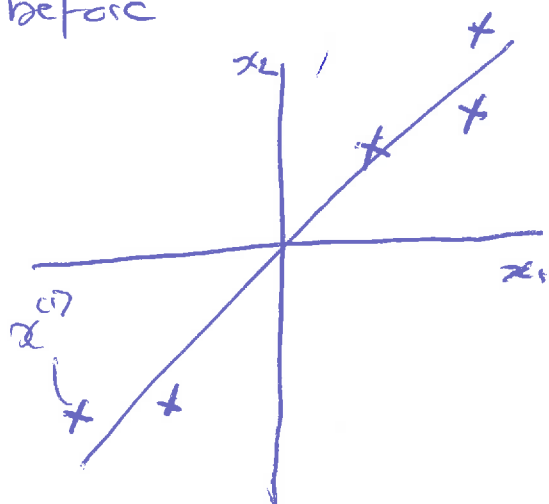
$$Z \in \mathbb{R} \rightarrow x \in \mathbb{R}^2$$

$$x_{\text{approx}} = \underbrace{U}_{n \times k} \cdot \underbrace{Z}_{k \times 1}$$

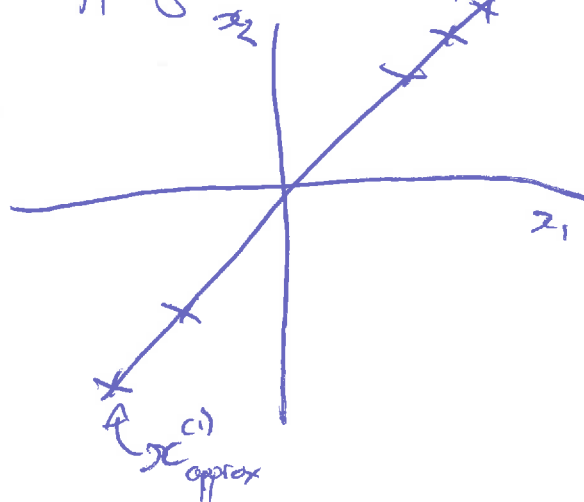
$\approx x$   
 $\uparrow$   
original  $x$

$n \times 1$

before



remapping back (after)



# Dimensionality Reduction - choosing the number of principal components

## Choosing $k$ (number of principal components)

Average squared projection error:  $\frac{1}{m} \sum_{i=1}^m \|x^{(i)} - x_{\text{approx}}^{(i)}\|^2$

Total variation in the data:  $\frac{1}{m} \sum_{i=1}^m \|x^{(i)}\|^2$

Typically, choose  $k$  to be smallest value so that

$$\frac{\frac{1}{m} \sum_{i=1}^m \|x^{(i)} - x_{\text{approx}}^{(i)}\|^2}{\frac{1}{m} \sum_{i=1}^m \|x^{(i)}\|^2} \leq 0.01 \quad (1\%)$$

or  
0.05 5%

"99% of variance is retained" / 95% of variance retained

## Choosing $k$ (number of principal components)

Algorithm:

Try PCA with  $k=1$ : if not less than 0.01, try  $k=2$ , if not try  $k=3$  etc

Compute Ureduce,  $z^{(1)}, z^{(2)}, \dots, z^{(m)}, x_{\text{approx}}^{(1)}, \dots, x_{\text{approx}}^{(m)}$

check if:

$$\frac{\frac{1}{m} \sum_{i=1}^m \|x^{(i)} - x_{\text{approx}}^{(i)}\|^2}{\frac{1}{m} \sum_{i=1}^m \|x^{(i)}\|^2} \leq 0.01$$

$$[U, S, V] = \text{svd}(\text{sigma}_A)$$

$$S = \begin{bmatrix} s_{11} & s_{22} & s_{33} & \dots & s_{nn} \\ 0 & & & & 0 \end{bmatrix}$$

For given value  $k$

$$1 - \frac{\sum_{i=1}^k s_{ii}}{\sum_{i=1}^n s_{ii}} \leq 0.01$$

$k=3$   
 $n=7$

## Summary

→ Pick smallest value of  $k$  for which

$$\frac{\sum_{i=1}^k s_{ii}}{\sum_{i=1}^n s_{ii}} \geq 0.99 \quad (99\% \text{ variance retained})$$

"choose what dimension to reduce the data too"

# Dimensionality reduction - advice for applying PCA

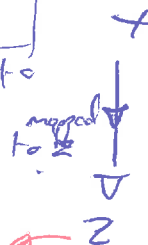
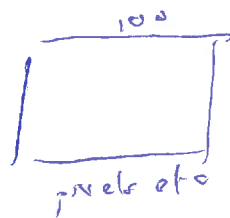
## Supervised learning speedup

$(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(n)}, y^{(n)})$

extract inputs:

Unlabeled dataset:  $x^{(1)}, x^{(2)}, \dots, x^{(n)} \in \mathbb{R}^{10000}$

$$x^{(i)} \in \mathbb{R}^{10000}$$



new representation

↓ PCA

$$z^{(1)}, z^{(2)}, \dots, z^{(n)} \in \mathbb{R}^{1000}$$

New training set

$(z^{(1)}, y^{(1)}), (z^{(2)}, y^{(2)}), \dots, (z^{(n)}, y^{(n)})$

if using logistic regression (for example):

$$h_{\theta}(z) = \frac{1}{1 + e^{-\theta^T z}}$$

Note: Mapping  $x^{(i)} \rightarrow z^{(i)}$  should be defined by running PCA only on the training set. This mapping can be applied as well to the examples  $x_{\text{dev}}^{(i)}$  and  $x_{\text{test}}^{(i)}$  in the cross validation and test sets.

Reduce

parameter learned by PCA

→ fit parameters only to training set. not cv or test set.

apply mapping too dev cv & test sets

## Application of PCA

- Compression
  - Reduce memory/disk needed to store data.
  - Speed up learning algorithm
  - choose  $k$  by % of variance retained.
- Visualization
  - $k=2$  or  $k=3$  ✓

## Dimensionality reduction - advice for applying PCA

Bad use of PCA: To prevent overfitting:

Use  $\underline{z}^{(i)}$  instead of  $x^{(i)}$  to reduce the number of features  $\overset{1000}{K} \leq \overset{1000}{n}$

Thus, fewer features, less likely to overfit. **Bad!**

This might work ok, but it is not a good way to address overfitting. Use regularization instead.

(PCR)  $\lambda$  ftw. ( $\lambda$  knows what the values of  $y$  are ... )  
minimization term  
→ it throws away data, without knowing what the values of  $y$  are.

PCA is sometimes used where it should not be.

Design of ML system:

→ Get training set  $((x^{(1)}, y^{(1)}) \dots (x^{(n)}, y^{(n)}))$

~~→ Run PCA to reduce  $x^{(i)}$  in dimension to get  $z^{(i)}$~~

→ train logistic regression on  $((z^{(1)}, y^{(1)}) \dots (z^{(n)}, y^{(n)}))$

→ test on test set: Map  $x_{\text{test}}^{(i)}$  to  $z_{\text{test}}^{(i)}$ . Run  $h_{\theta}(z)$  on  
 $(z_{\text{test}}^{(1)}, y_{\text{test}}^{(1)}) \dots (z_{\text{test}}^{(n)}, y_{\text{test}}^{(n)})$

→ How about the whole thing without PCA

→ try running the normal thing with original / raw data  $x^{(i)}$ . If the implementation is too slow etc try implementing  $z^{(i)}$  etc