

Dimension Reduction 2: PCA

Feb 17 2022

J.L.:

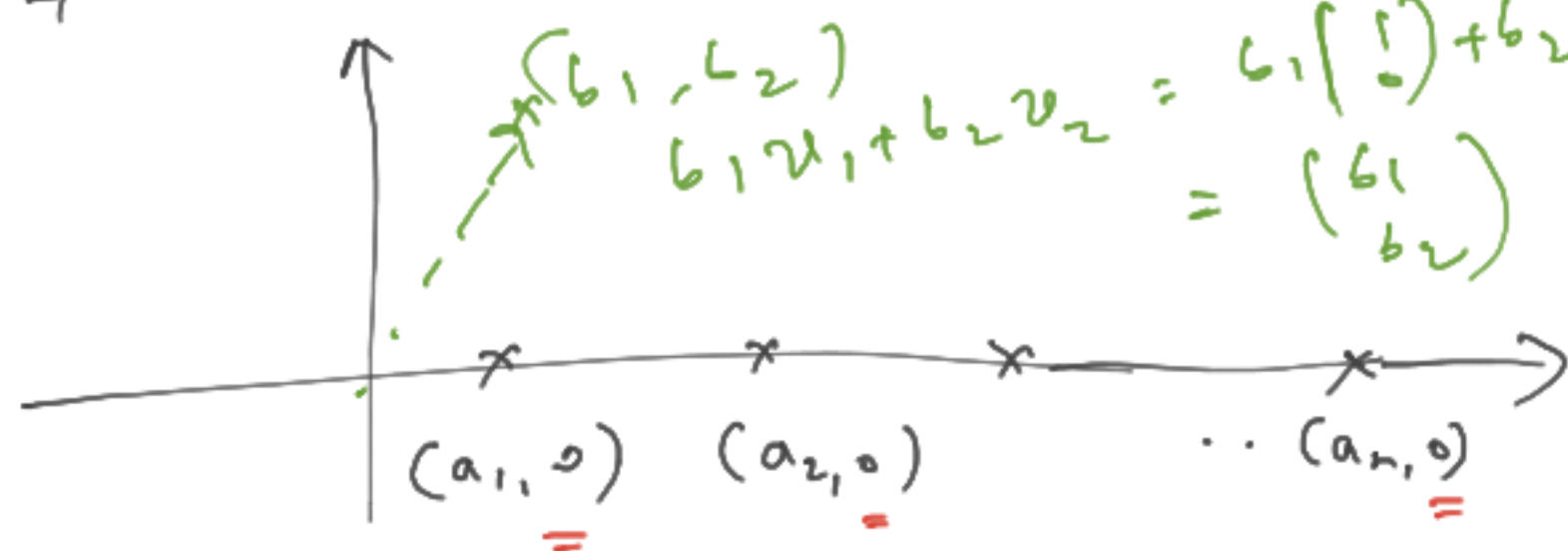
+ve: data independent

-ve: ——— " ———
cannot exploit structure
in data.

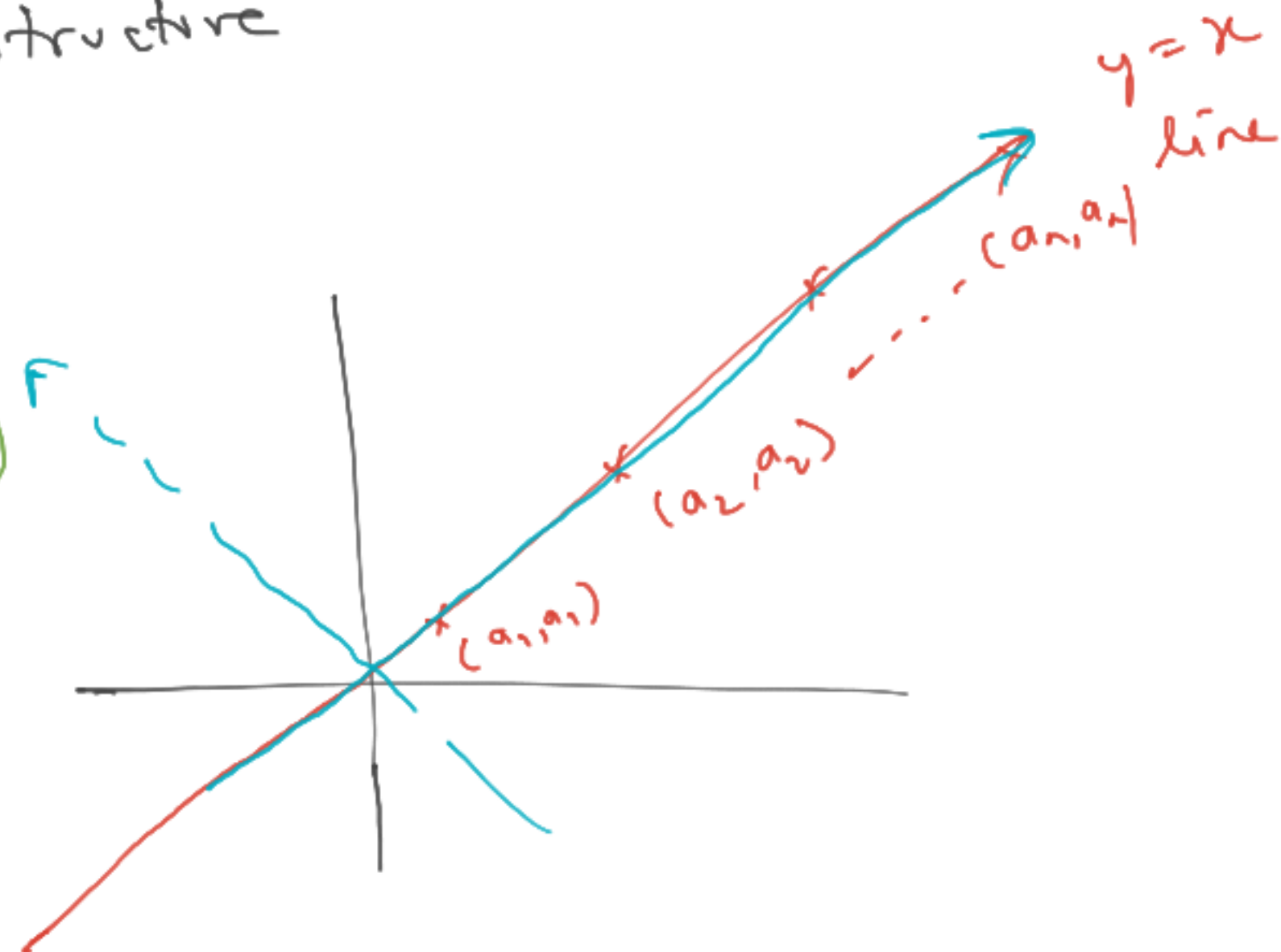
$$K = O\left(\frac{\log n}{\epsilon^2}\right)$$

indep. of D

Toy example:

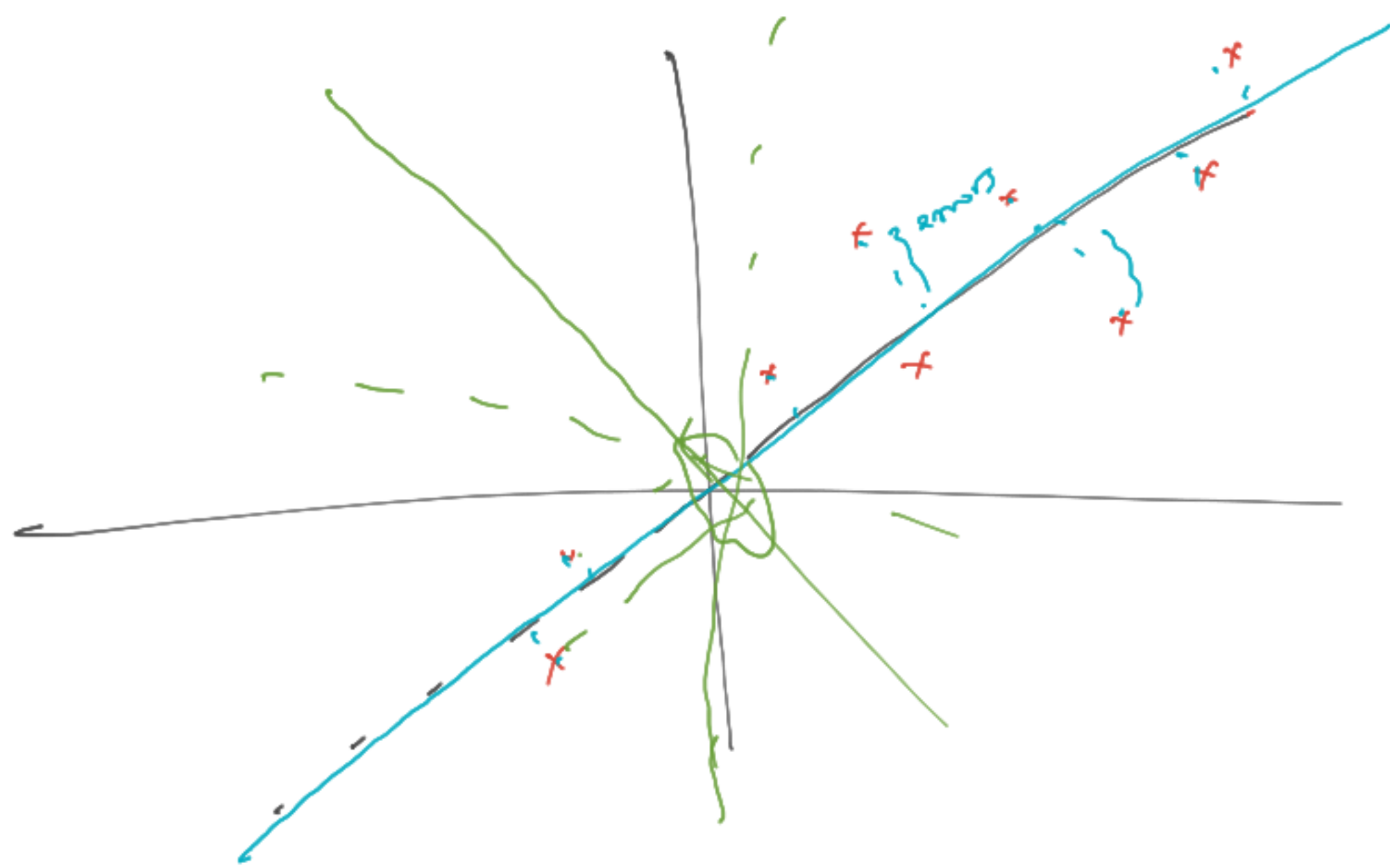


$$(b_1, b_2)$$



$$v_1 = (1, 0) \rightarrow x\text{-axis}$$

$$v_2 = (0, 1) \rightarrow y\text{-axis}$$



PCA/SVD:

Maximize variance/info
in the "chosen"
dimensions.

Applications:

1. Analysis of genome data: Gene microarray data
original dimension $D = 10,000$ genes
 n vectors in \mathbb{R}^D
↓
(individuals)
2. Denoising stock market data

Principal Component Analysis (PCA)

Goal of PCA: To find K orthonormal vectors s.t. the points in the dataset have a good approximation in the subspace generated by these vectors

Real
'basis
vectors'

Good approximation: is in L_2 sense

↓

(squared error)

between original points and their approximation
in the low-dimensional subspace

$\{u_1, u_2, \dots, u_k\}$ orthonormal "basis" vectors

Looking at subspace spanned by these vectors

$$\hat{\omega} = \sum_{i=1}^k \alpha_i u_i \in \mathbb{R}^D$$

\downarrow
 scalars

$$(\alpha_1, \alpha_2, \dots, \alpha_k) \in \mathbb{R}^k$$

$$u_i \in \mathbb{R}^D$$

$$\omega \in \mathbb{R}^D$$

arbitrary vector

Approximation of vector ω in the chosen subspace

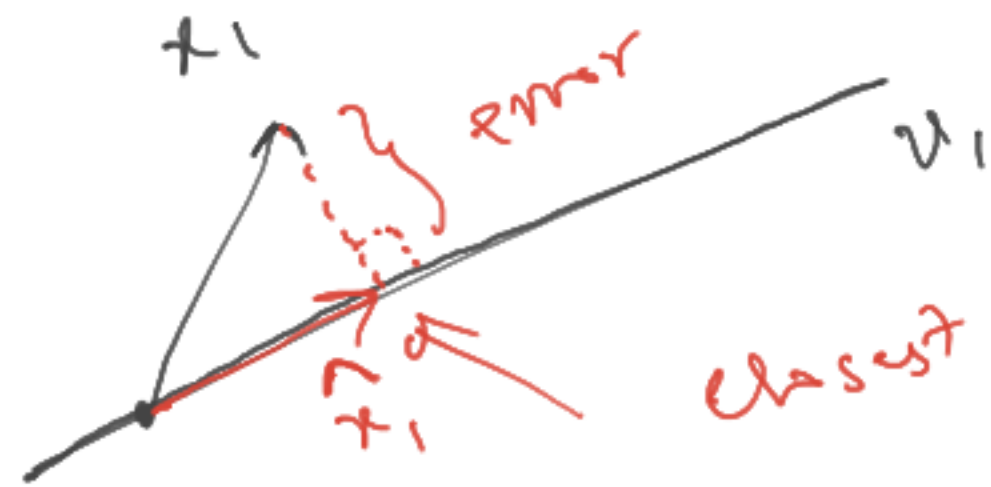
$$\hat{\omega} = \alpha_1 u_1 + \alpha_2 u_2 + \dots + \alpha_k u_k \in \mathbb{R}^D$$

Error $\|\omega - \hat{\omega}\|$

$$\begin{aligned} \hat{x}_1 &= (\text{scalar}) u_1 \\ &= \langle x_1, u_1 \rangle u_1 \end{aligned}$$

\uparrow
inner product

$$= (x_1^T u_1) u_1$$



closest approx of x_1 along u_1 = projection of x_1 along u_1

By Pythagoras,

$$\|x_p\|^2 = \underbrace{|\text{projection}|^2}_{\uparrow \text{maximize}} + \underbrace{|\text{error}|^2}_{\downarrow \text{minimize}}$$

\therefore Minimizing l_2 error of approx.
= maximizing projected distances
= maximizing the variance/information in
the chosen dimensions

Preprocessing:

PCA is very sensitive to scaling

⇒ Need preprocessing.

1. data zero mean (by subtracting the sample mean)
2. Scale each co-ordinate so that they are comparable
(divide by sample deviation)

Solving PCA : 1-dim.

Given a unit vector u
a point x

} length of the projection
of x onto u
in $x^T u$.

n data points x_1, \dots, x_n

Find a vector u s.t maximize the projected distances.

$$\begin{aligned} \text{maximize} \quad & \sum_{i=1}^n (x_i^T u)^2 = \sum_{i=1}^n u^T x_i x_i^T u \\ & = u^T \left(\sum_{i=1}^n x_i x_i^T \right) u \\ & = \text{maximize} \quad u^T \underline{\underline{M}} u \\ & \quad \rightarrow \text{Covariance Matrix} \end{aligned}$$

Note: A blue arrow points from the term $x_i x_i^T$ in the second line to a diagram of a vector x_i and its projection $x_i^T u$ onto a line defined by vector u .

$$\arg \max_{\substack{u \in \mathbb{R}^D \\ \|u\|_2 = 1}} u^T M u = \text{principal eigen vector of } M$$

K-dimensions:

Let $M =$

$$\sum_{i=1}^n$$

$$x_i x_i^T = X^T X$$

→ Covariance matrix

Pos. Sem. Def (PSD)

⇒ Non-neg. eigen values

where

$$X = \begin{bmatrix} - & x_1^T & - \\ - & x_2^T & - \\ & \vdots & \\ - & x_n^T & - \end{bmatrix}$$

Eigen value Decomposition (EVD) of $M = V \Lambda V^T$

=

$$\begin{bmatrix} \lambda_1 & & 0 & \dots \\ & \lambda_2 & & \\ & & \ddots & \\ 0 & & & \dots \end{bmatrix}$$

↗ Eigen values
 ↘ Eigen vectors

Let $\lambda_1 \geq \lambda_2 \geq \lambda_3 \dots$

Let u_1, u_2, \dots, u_k be the first k columns of V .

Then $\{u_1, \dots, u_k\}$ are k orthonormal vectors that

maximize the projected distances (i.e. variance)

Transformation matrix

$S =$

$$\begin{bmatrix} u_1^T \\ u_2^T \\ \vdots \\ u_k^T \end{bmatrix}$$

Computing eigen values/vectors

Power Iteration:

High level steps:

1. Compute the principal eigen vector

2. Subtract out its contribution

3. Iterate

1. Pick a random vector y_0

2.
$$y_{i+1} = \frac{M y_i}{\|M y_i\|_2}$$

rescaling to norm 1

3. Iterate until convergence

to get (approximate) u_1

4. $\lambda_1 = v_1^T M v_1$

→ (lin. algebra concept)

$$(\text{mean})_M = \sum_{i=1}^n x_i x_i^T$$

$$M - \lambda, u, v, T$$

Intuition behind power iteration:

$$\text{Let } y_0 = \sum_{i=1}^K \alpha_i v_i = \alpha_1 v_1 + \alpha_2 v_2 + \dots + \alpha_k v_k$$

$$y_1 = M y_0 = (V \Lambda V^T) (\alpha_1 v_1 + \dots + \alpha_k v_k)$$

$$M = V \Lambda V^T$$

↓

orthonormal
columns

'diagonal'

$$= [\alpha_1 \lambda_1 v_1 + \alpha_2 \lambda_2 v_2 + \dots + \alpha_k \lambda_k v_k]$$

$$y_t = [\alpha_1 \lambda_1^t v_1 + \alpha_2 \lambda_2^t v_2 + \dots + \alpha_k \lambda_k^t v_k]$$

Projection of y_t along v_1

$$|\langle y_t, v_1 \rangle| = \frac{\alpha_1 \lambda_1^t}{\sqrt{\sum_{i=1}^K (\alpha_i \lambda_i^t)^2}}$$

↳ goes down to λ_1 / λ_2

When PCA does not work?

1. find "linear approximation" :

$$y = x^2$$

⇒ if there are nonlinear relationships

2. ↗ normalization is not done correctly

Exercise: Recall relationship btwn PCA and SVD