

CPSC 340: Machine Learning and Data Mining

More PCA
Spring 2022 (2021W2)

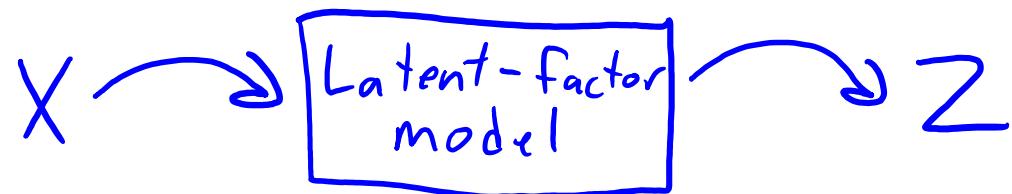
1. Decision trees
2. Naïve Bayes classification
3. Ordinary least squares regression
4. Logistic regression
5. Support vector machines
6. Ensemble methods
7. Clustering algorithms
8. Principal component analysis
9. Singular value decomposition
10. Independent component analysis (bonus)

The 10 Algorithms Machine Learning Engineers Need to Know

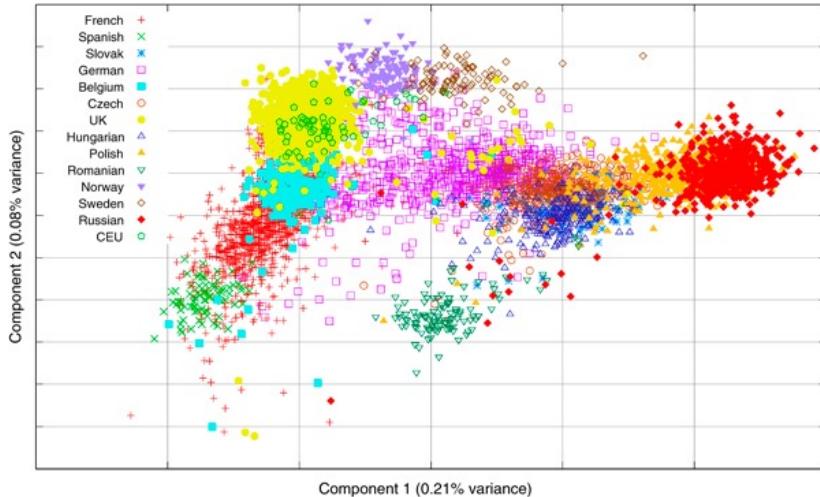


Last Time: Latent-Factor Models

- Latent-factor models take input data 'X' and output a basis 'Z':



- Usually, 'Z' has fewer features than 'X'.
- Uses: dimensionality reduction, visualization, factor discovery.



Trait	Description
Openness	Being curious, original, intellectual, creative, and open to new ideas.
Conscientiousness	Being organized, systematic, punctual, achievement-oriented, and dependable.
Extraversion	Being outgoing, talkative, sociable, and enjoying social situations.
Agreeableness	Being affable, tolerant, sensitive, trusting, kind, and warm.
Neuroticism	Being anxious, irritable, temperamental, and moody.

<http://infoproc.blogspot.ca/2008/11/european-genetic-substructure.html>

<https://new.edu/resources/big-5-personality-trait>

Last Time: Principal Component Analysis

- Principal component analysis (PCA) is a linear latent-factor model:
 - These models “factorize” matrix X into matrices Z and W :

$$X \approx ZW$$

$n \times d$

$$x_i \approx W^T z_i$$

$$x_{ij} \approx \langle w_j^T, z_i \rangle$$

- We can think of rows w_c of W as ‘ k ’ fixed “part” (used in all examples).
- z_i is the “part weights” for example x_i : “how much of each part w_c to use”.

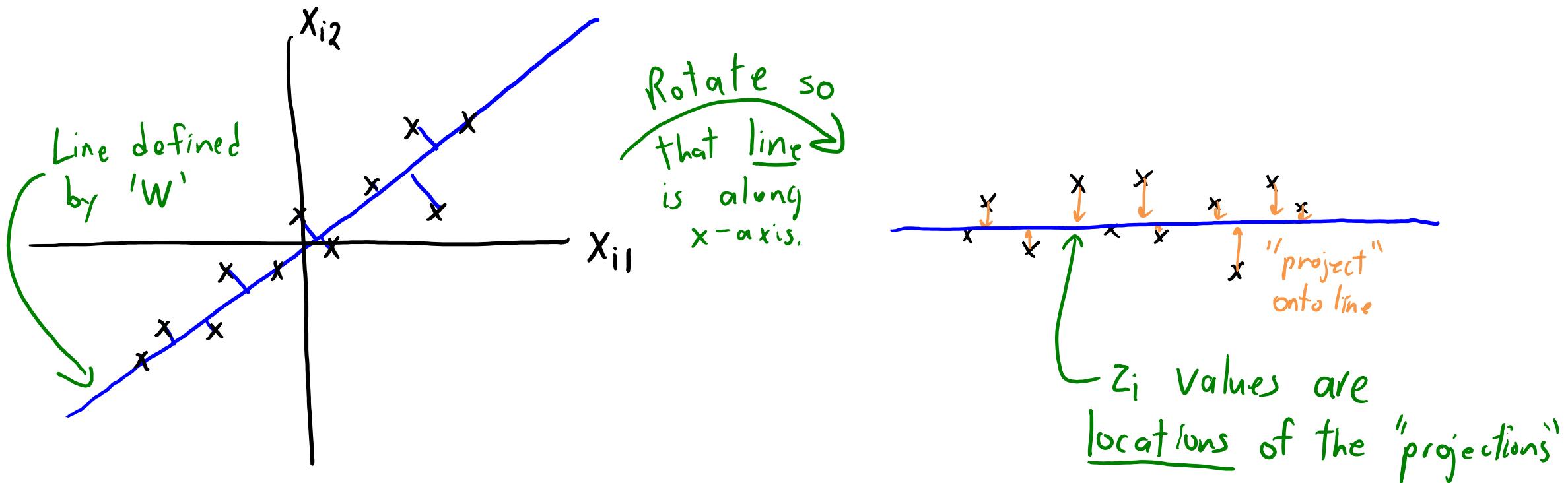
$$\begin{matrix} 5 \\ = 1 & + 0 & + 1 & + 1 & + 1 & + 0 & + 1 \end{matrix}$$

$\hat{x}_i = z_{i1} w_1 + z_{i2} w_2 + z_{i3} w_3 + z_{i4} w_4 + z_{i5} w_5 + z_{i6} w_6 + z_{i7} w_7$

"parts" are called "principal components" or "factors"

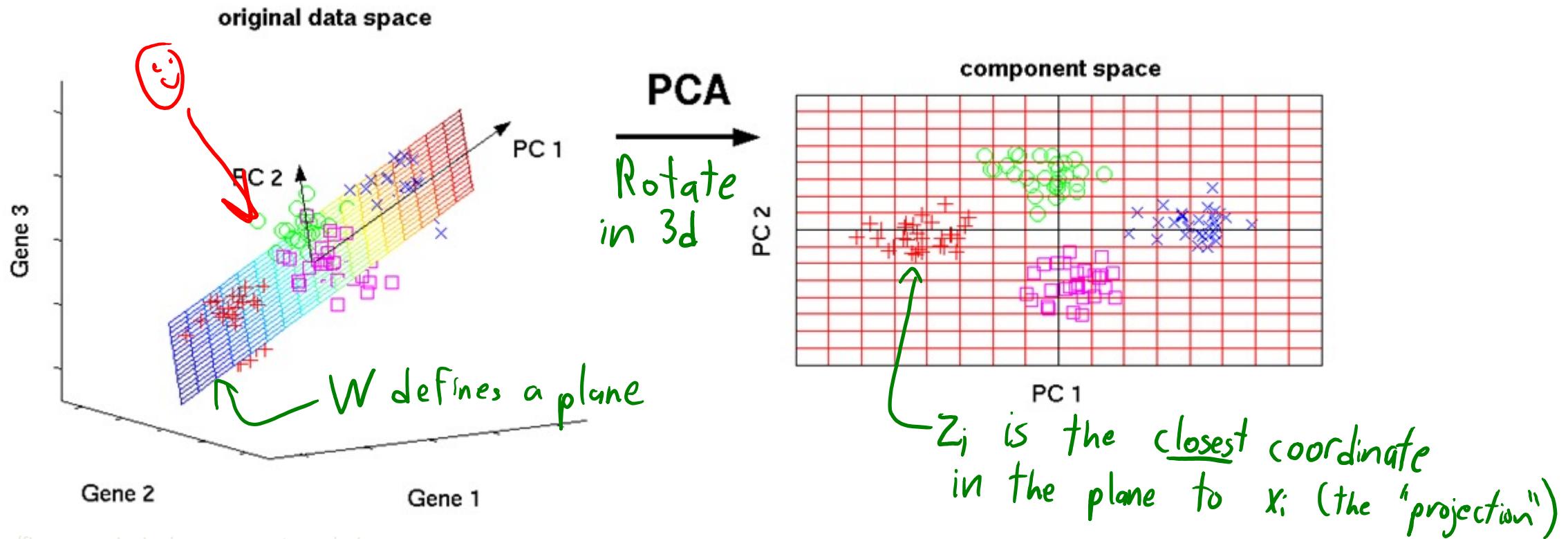
Last Time: PCA Geometry

- When $k=1$, the W matrix defines a line:
 - We choose ' W ' as the line minimizing squared distance to the data.
 - Given ' W ', the z_i are the coordinates of the x_i “projected” onto the line.



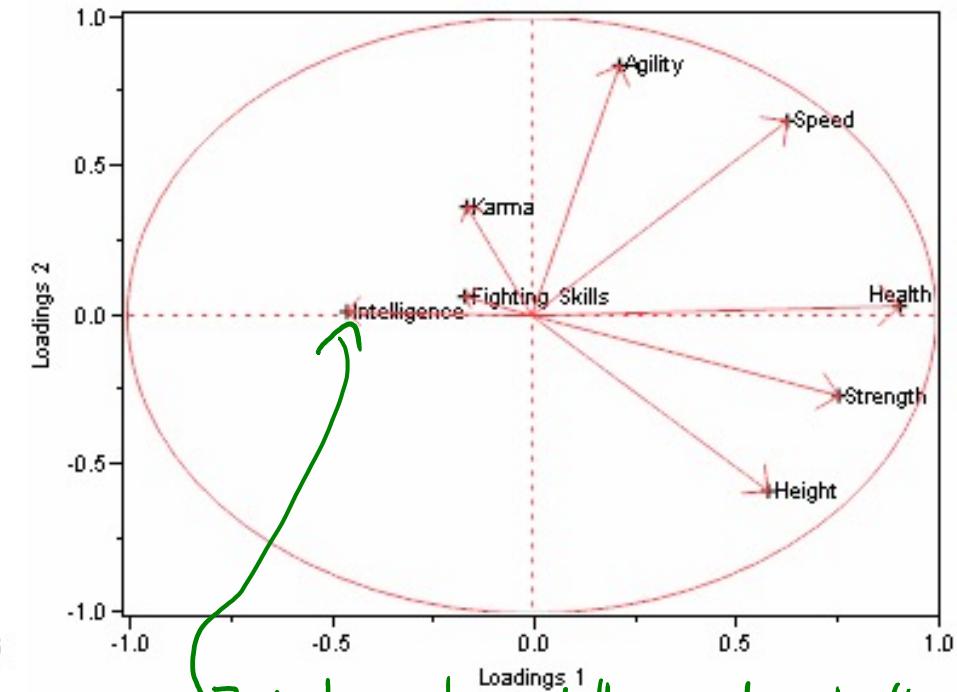
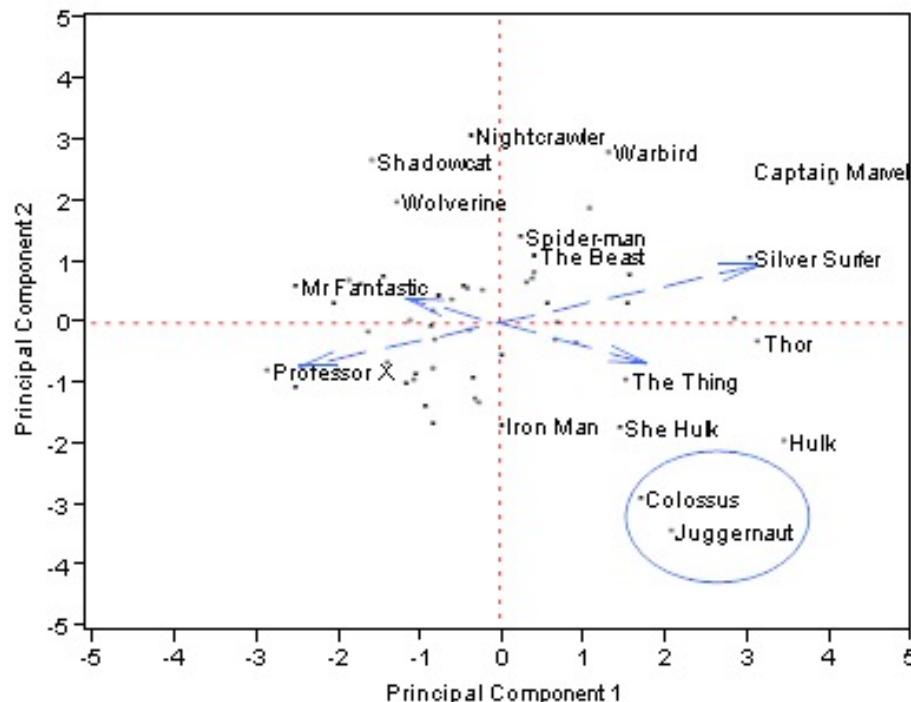
Last Time: PCA Geometry

- When $k=2$, the W matrix defines a plane:
 - We choose ‘ W ’ as the plane minimizing squared distance to the data.
 - Given ‘ W ’, the z_i are the coordinates of the x_i “projected” onto the plane.



Last Time: PCA Geometry

- When $k=2$, the W matrix defines a **plane**:
 - Even if the original data is high-dimensional,
we can **visualize** data “projected” onto this plane.



PCA Objective Function

- In PCA we minimize the squared error of the approximation:

$$f(W, z) = \sum_{i=1}^n \|W^T z_i - \underbrace{x_i}_{\text{approximation example } i}\|^2$$

- This is equivalent to the k-means objective:
 - In k-means z_i only has a single '1' value and other entries are zero.
- But in PCA, z_i can be any real number.
 - We approximate x_i as a linear combination of all means/factors.

PCA Objective Function

- In PCA we minimize the squared error of the approximation:

$$f(W, z) = \sum_{i=1}^n \|W^T z_i - x_i\|^2 = \sum_{i=1}^n \sum_{j=1}^d (\langle w_j^i, z_i \rangle - x_{ij})^2$$

approximation example 'i' approximating feature 'j' of example 'i'

- We can also view this as solving 'd' regression problems:
 - Each w_j^i is trying to predict column ' x_j ' from the basis z_i .
 - The output " y_i " we try to predict here is actually the features " x_i ".
 - But unlike in regression, we're also learning the features z_i .

Principal Component Analysis (PCA)

- The 3 different ways to write the PCA objective function:

$$\begin{aligned} f(W, Z) &= \sum_{i=1}^n \sum_{j=1}^d (\langle w_j^T z_i \rangle - x_{ij})^2 && \text{(approximating } x_{ij} \text{ by } \langle w_j^T z_i \rangle \text{)} \\ &= \sum_{i=1}^n \|W^T z_i - x_i\|^2 && \text{(approximating } x_i \text{ by } W^T z_i \text{)} \\ &= \|Z^T W - X\|_F^2 && \text{(approximating } X \text{ by } Z^T W \text{)} \end{aligned}$$

Digression: Data Centering (Important)

- In PCA, we assume that the data X is “centered”.
 - Each column of X has a mean of zero.
- It's easy to center the data:

Set $\mu_j = \frac{1}{n} \sum_{i=1}^n x_{ij}$ (mean of column ' j ')

Replace each x_{ij} with $(x_{ij} - \mu_j)$

- There are PCA variations that estimate “bias in each coordinate”.
 - In basic model this is equivalent to centering the data.

PCA Computation: Prediction

- At the end of training, the “model” is the μ_j and the W matrix.
 - PCA is parametric.
- PCA prediction phase:
 - Given new data \tilde{X} , we can use μ_j and W this to form \tilde{Z} :

1. Center: replace each \tilde{x}_{ij} with $(\tilde{x}_{ij} - \mu_j)$

2. Find \tilde{Z} minimizing squared error:

$$\tilde{Z} = \tilde{X} W^T (W W^T)^{-1}$$

(could just store
this $d \times k$ matrix)

$\overbrace{\mu_j}$ means of
training
data

PCA Computation: Prediction

- At the end of training, the “model” is the μ_j and the W matrix.
 - PCA is parametric.
- PCA prediction phase:
 - Given new data \tilde{X} , we can use μ_j and W this to form \tilde{Z} :
 - The “reconstruction error” is how close approximation is to \tilde{X} :

$$\|\underbrace{\tilde{Z}W}_{\hat{X}} - \underbrace{\tilde{X}}_{\text{Centered version}}\|_F^2$$

- Our “error” from replacing the x_i with the z_i and W .

Choosing ‘k’ by “Variance Explained”

- Common to choose ‘k’ based on variance of the x_{ij} .

$$\text{Var}(x_{ij}) = E[(x_{ij} - \mu_{ij})^2] = E[x_{ij}^2] = \frac{1}{nd} \sum_{i=1}^n \sum_{j=1}^d x_{ij}^2 = \frac{1}{nd} \|X\|_F^2$$

definition of variance
assumed to be zero
definition of expectation
Frobenius norm

- For a given ‘k’ we compute (variance of errors)/(variance of x_{ij}):

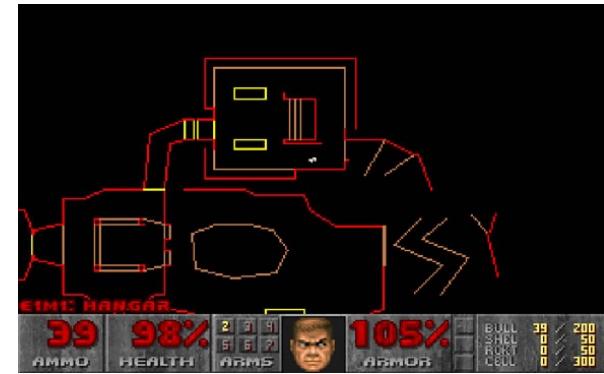
$$\frac{\|ZW - X\|_F^2}{\|X\|_F^2}$$

Centred version

- Gives a number between 0 ($k=d$) and 1 ($k=0$), giving “variance remaining”.
 - If you want to “explain 90% of variance”, choose smallest ‘k’ where ratio is < 0.10.

“Variance Explained” in the Doom Map

- Recall the Doom latent-factor model (where map ignores height):



- Interpretation of “variance remaining” formula:

$$\frac{\|Z_W - X\|_F^2}{\|X\|_F^2} \leftarrow \begin{array}{l} \text{Variance in } z\text{-dimension (variance in } x\text{- and } y\text{-dimensions fully} \\ \text{captured by overhead map)} \end{array}$$

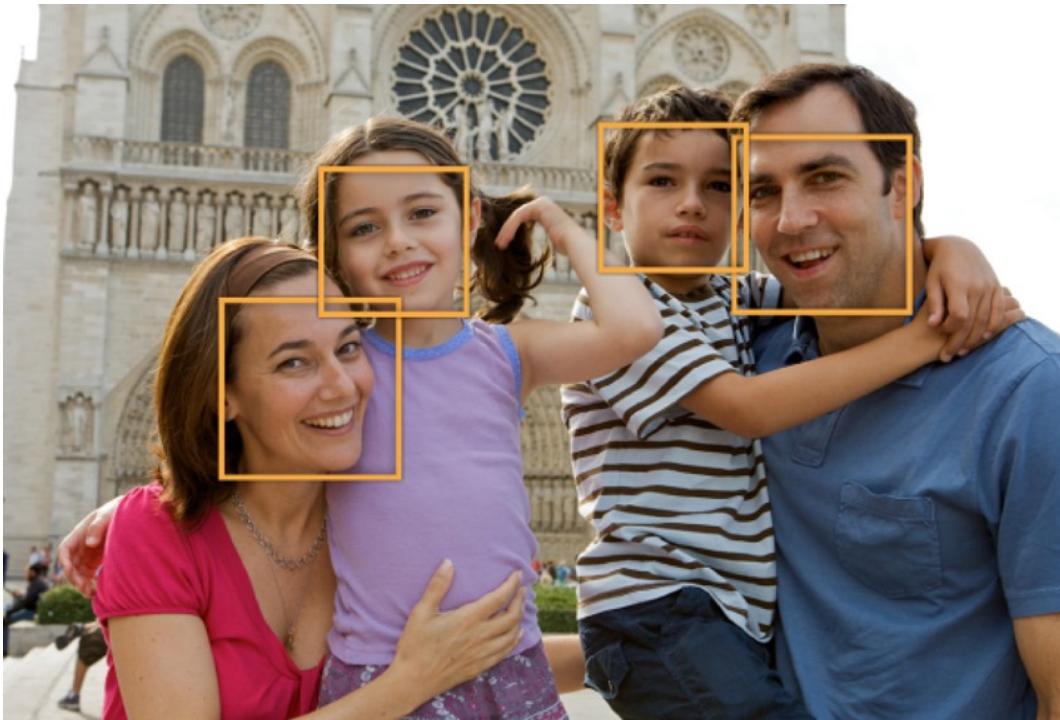
$\|X\|_F^2 \leftarrow \text{Variance of character in 3-dimensions}$

- If we had a 3D map the “variance remaining” would be 0.

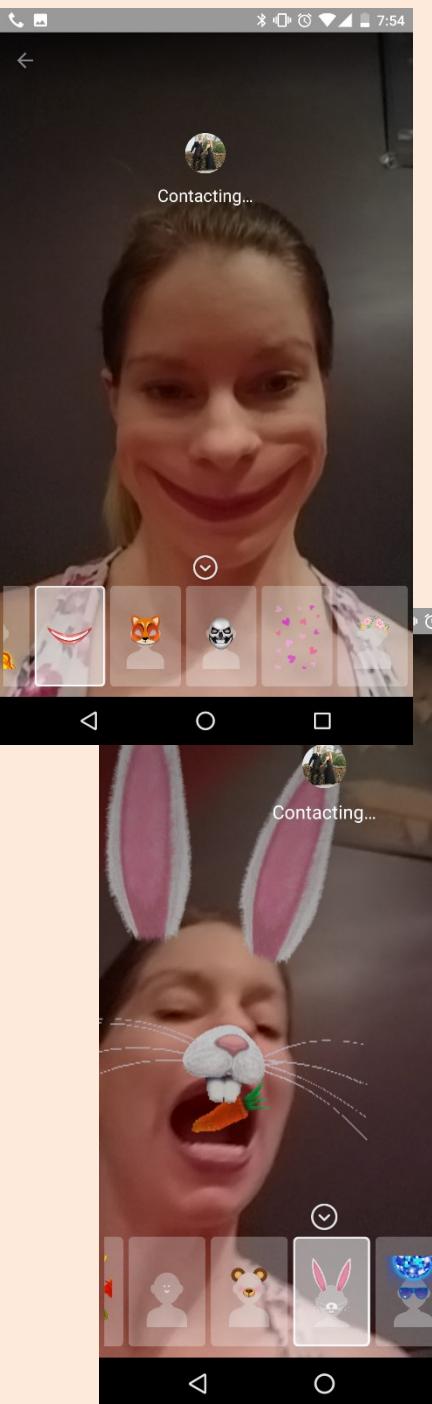
(pause)

Application: Face Detection

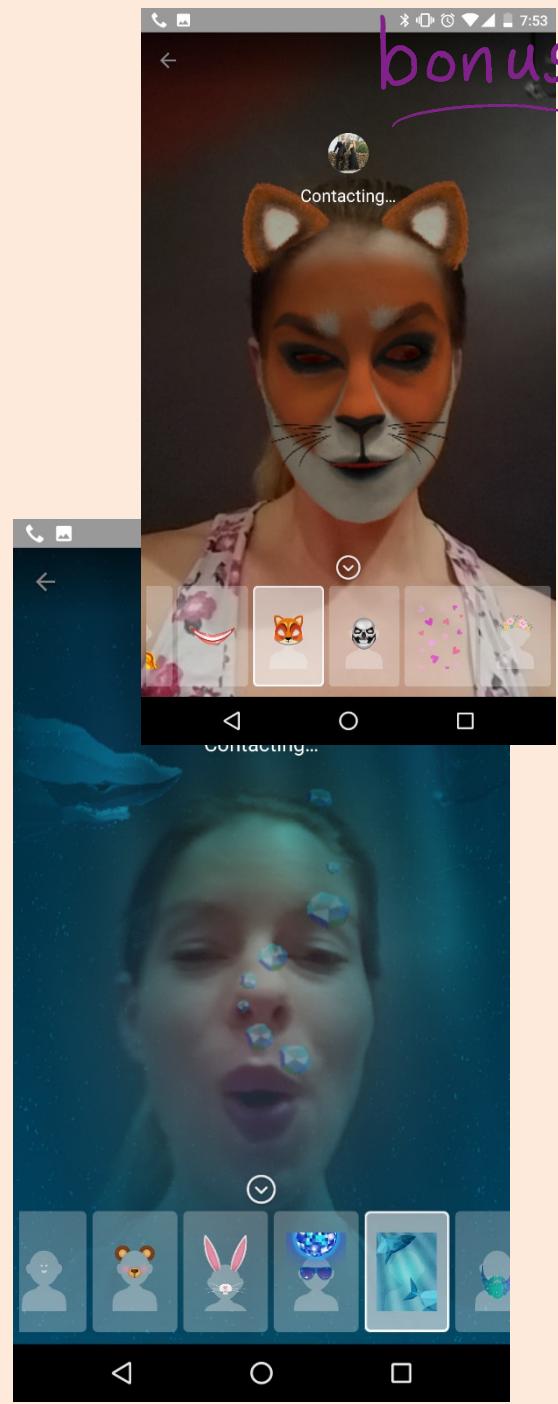
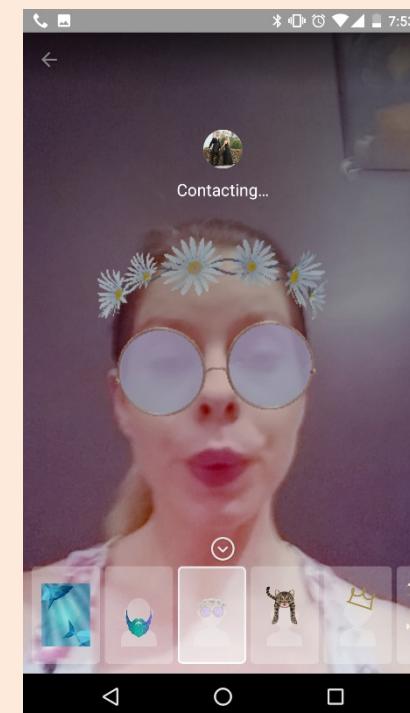
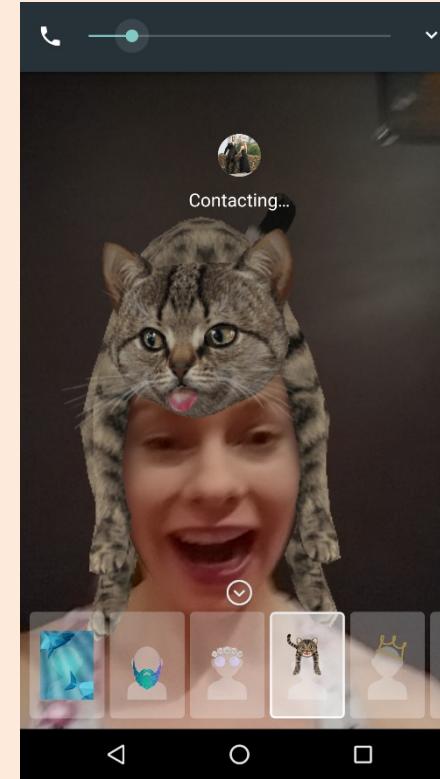
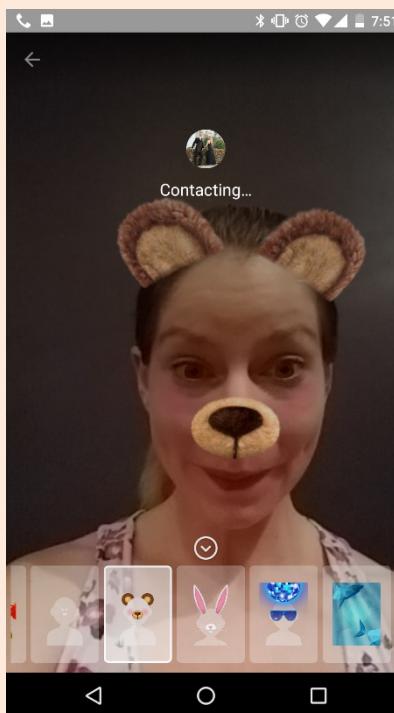
- Consider problem of face detection:



- Classic methods use “eigenfaces” as basis:
 - PCA applied to images of faces.



Application: Face Detection



Eigenfaces

- Collect a bunch of images of faces under different conditions:



Each row of X will be pixels in one image:

$$X =$$

If have ' n ' images that are ' m ' by ' m ' then X is ' n ' by m^2 .

Eigenfaces

Compute mean μ_j of each column.



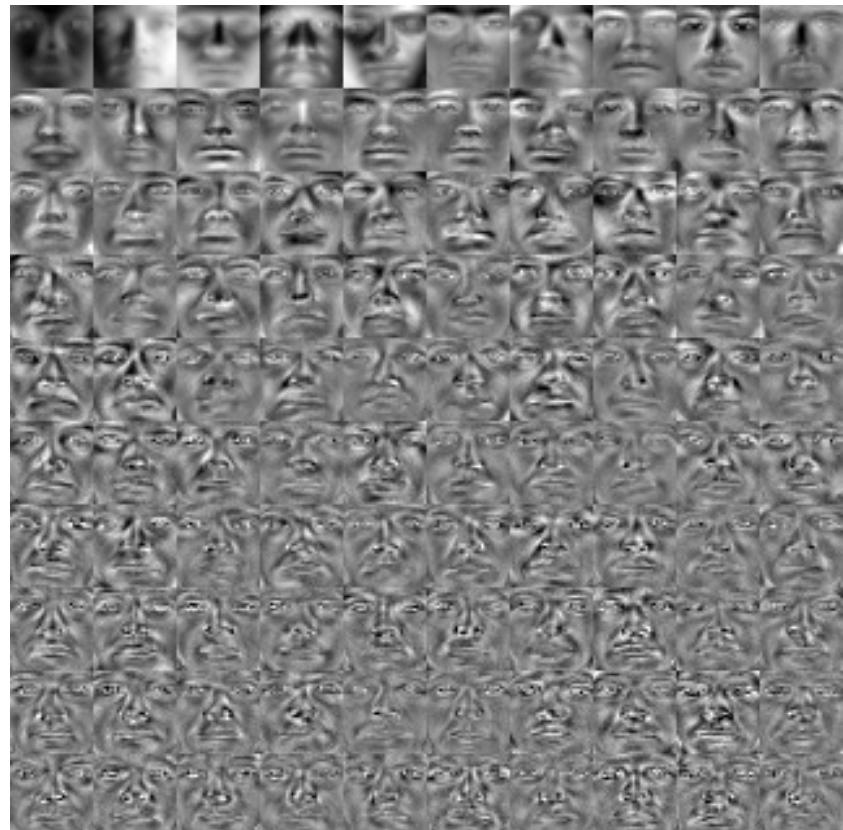
Each row of X will be pixels in one image:

$$X = \begin{bmatrix} x_1 - \mu \\ x_2 - \mu \\ \vdots \\ x_n - \mu \end{bmatrix}$$

Replace each x_{ij} by $x_{ij} - \mu_j$

Eigenfaces

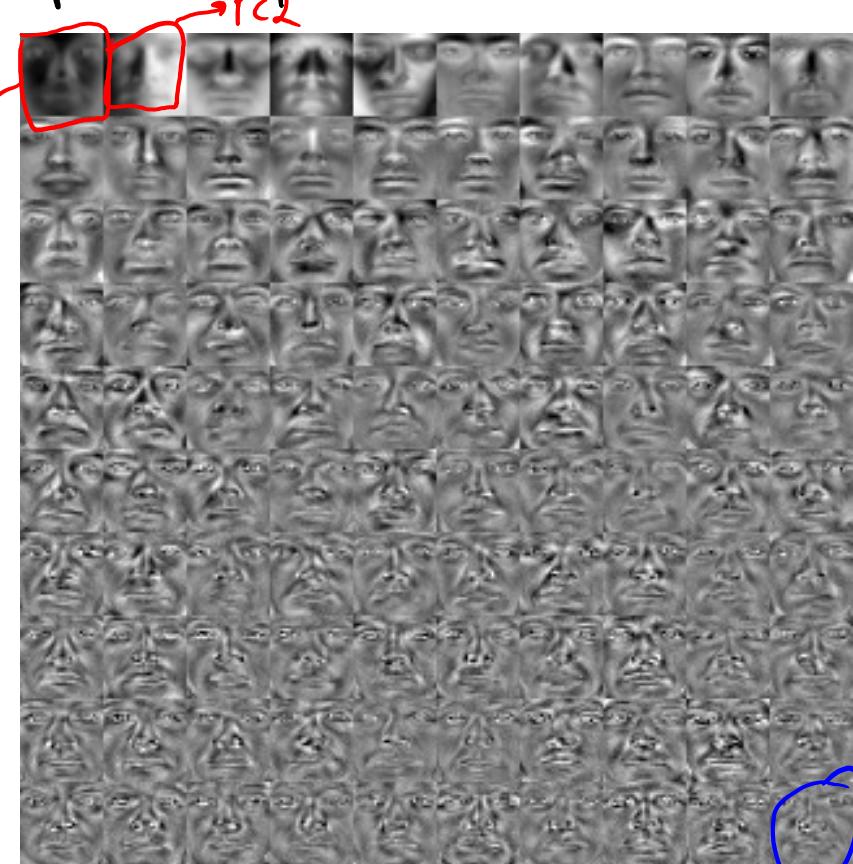
Compute top 'k' PCs on centered data: Each row of X will be pixels in one image:



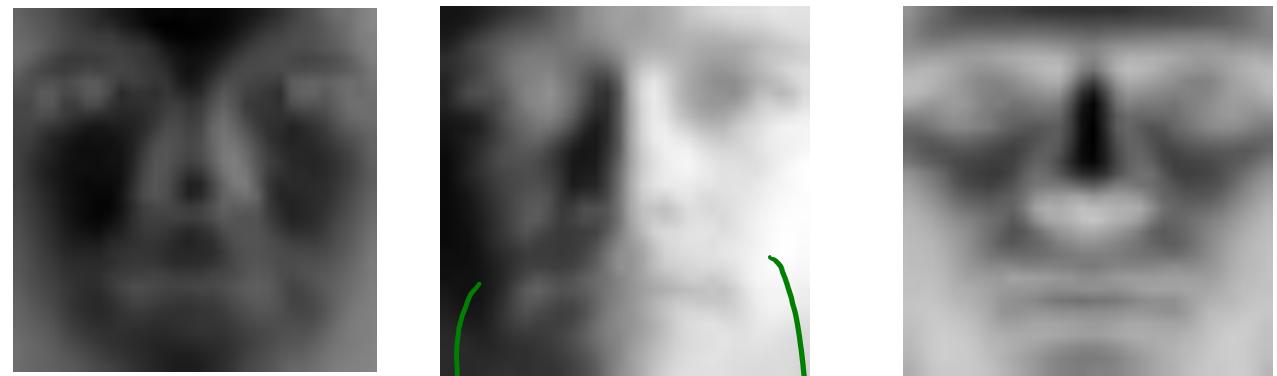
$$X = \begin{bmatrix} x_1 - \mu \\ x_2 - \mu \\ \vdots \\ x_n - \mu \end{bmatrix}$$

Eigenfaces

Compute top 'k' PCs on centered data:



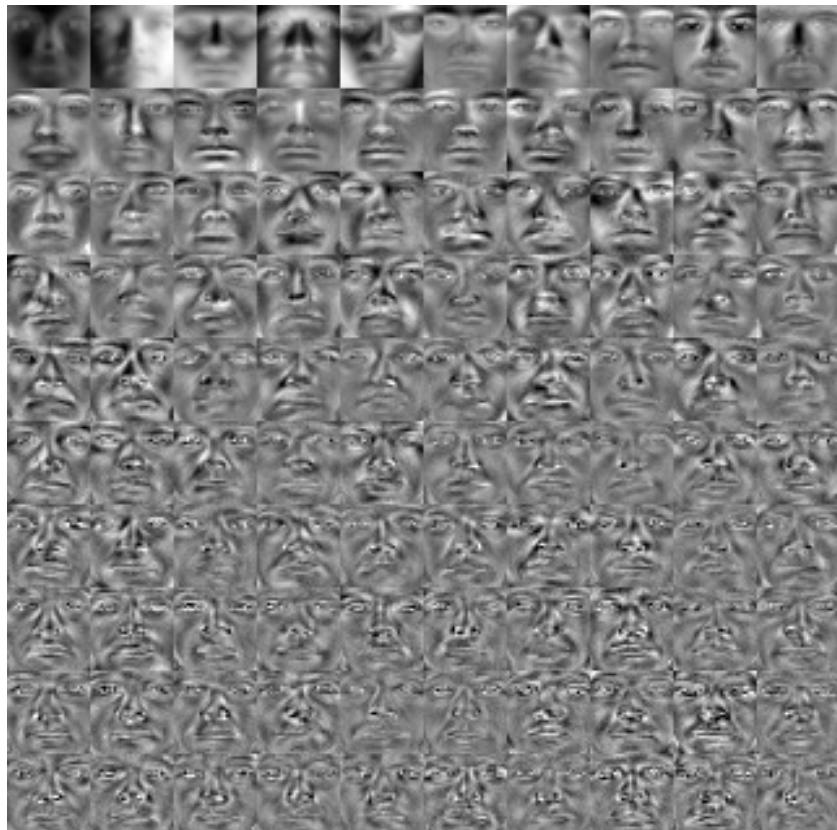
Note that these are "signed" images.



"gray" represents values close to 0.
"dark" represents negative values
"bright" represents positive values

Eigenfaces

Compute top 'k' PCs on centered data:



"Eigenface" representation:

$$\hat{x}_i = \mu + z_{i1} \text{PC1} + z_{i2} \text{PC2} + z_{i3} \text{PC3} + \dots$$

\hat{x}_i μ z_{i1} PC1
 $(\text{first row of } W)$ z_{i2} PC2 z_{i3} PC3 \dots

Eigenfaces

106 of the original faces:



"Eigenface" representation:

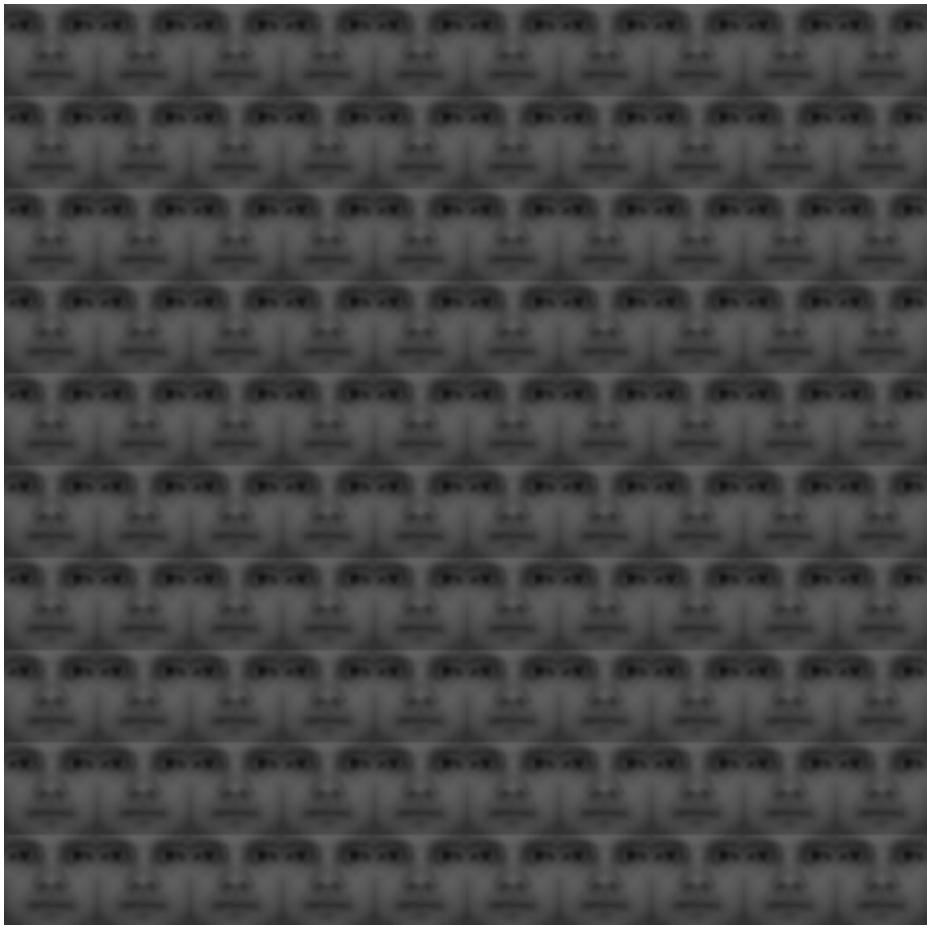
$$\hat{x}_i = \mu + z_{i1} \text{PC1} + z_{i2} \text{PC2} + z_{i3} \text{PC3} + \dots$$

\hat{x}_i μ z_{i1} PC1
 z_{i2} PC2
 z_{i3} PC3
 \vdots

(first row of W)

Eigenfaces

Reconstruction with $k=0$



Variance explained: 0%

"Eigenface" representation:

$$\hat{x}_i = \mu + z_{i1} \text{PC1} + z_{i2} \text{PC2} + z_{i3} \text{PC3} + \dots$$

\hat{x}_i μ z_{i1} z_{i2} z_{i3} PC1 PC2 PC3

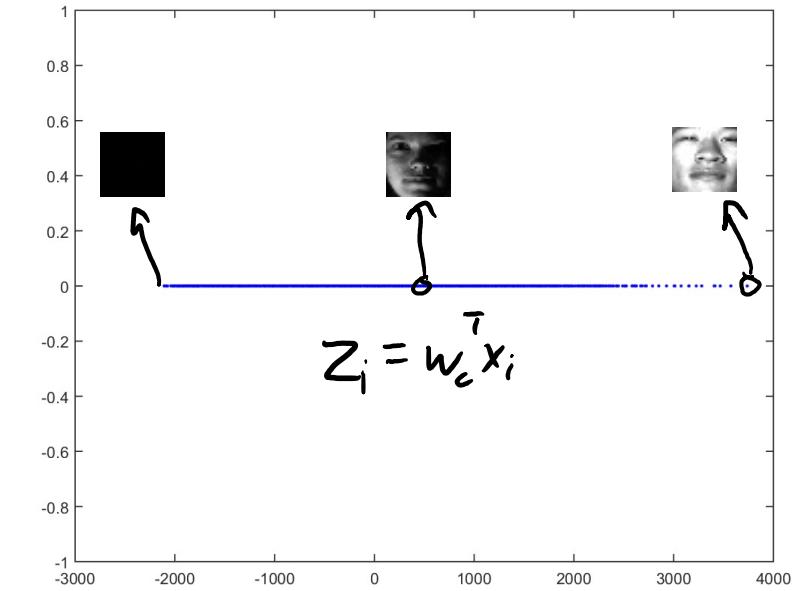
(first row of W)

Eigenfaces

Reconstruction with $k=1$



PCA Visualization:



"Eigenface" representation:

Variance explained: 36%

$$\hat{x}_i = \mu + z_{i1} \text{PC1} + z_{i2} \text{PC2} + z_{i3} \text{PC3} + \dots$$

The diagram shows the decomposition of a face \hat{x}_i into its mean μ and components along principal components PC1, PC2, PC3, etc. A red box highlights the first term $\hat{x}_i = \mu + z_{i1} \text{PC1}$, with a red bracket underneath labeled "(first row of W)".

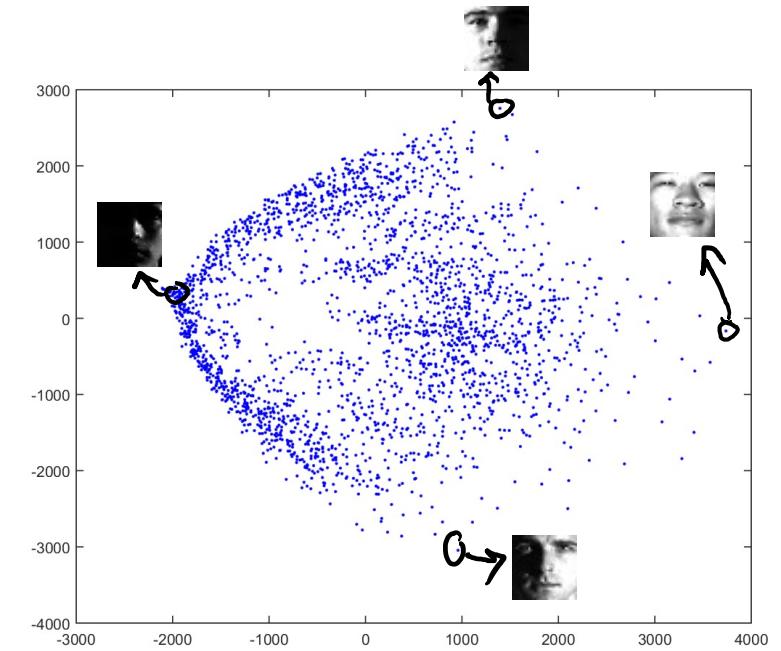
Eigenfaces

Reconstruction with $k=2$



Variance explained: 71%

PCA Visualization:



"Eigenface" representation:

$$\hat{x}_i = \mu + z_{i1} \text{PC1} + z_{i2} \text{PC2} + z_{i3} \text{PC3} + \dots$$

The equation shows the "Eigenface" representation of a face \hat{x}_i . It is represented as the sum of the mean face μ , the first principal component PC1 scaled by the coefficient z_{i1} , the second principal component PC2 scaled by z_{i2} , the third principal component PC3 scaled by z_{i3} , and so on. A red oval encloses the terms \hat{x}_i , μ , PC1 , PC2 , PC3 , and the ellipsis. Below the PC terms, it says "(first row of W)".

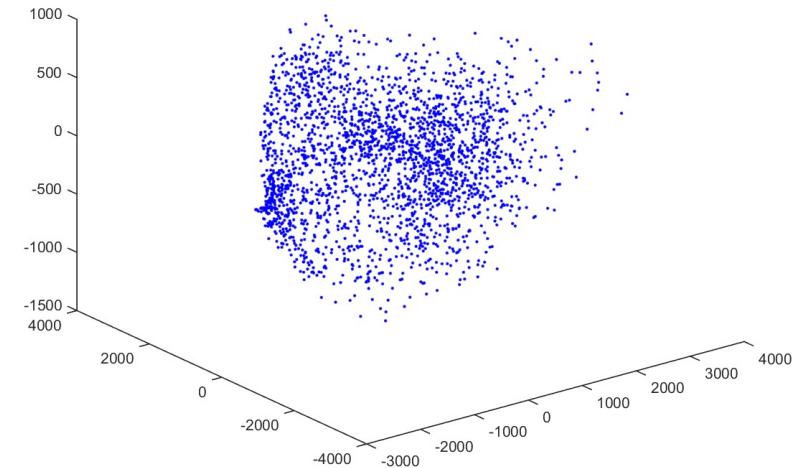
Eigenfaces

Reconstruction with $k=3$



Variance explained: 76%

PCA Visualization:



"Eigenface" representation:

$$\hat{x}_i = \mu + z_{i1} \text{PC1} + z_{i2} \text{PC2} + z_{i3} \text{PC3} + \dots$$

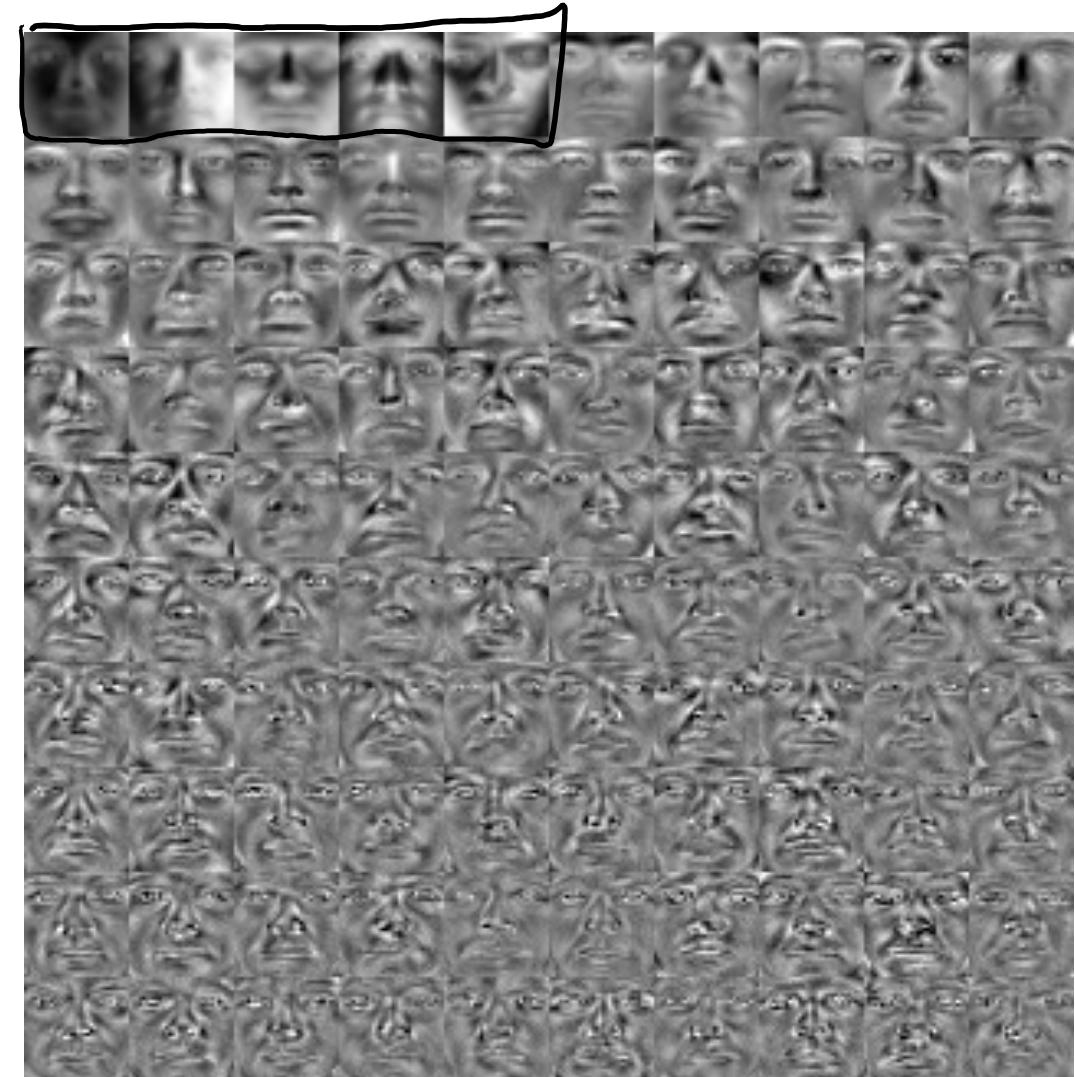
(first row of W)

Eigenfaces

Reconstruction with $k=5$



Variance explained: 80 %

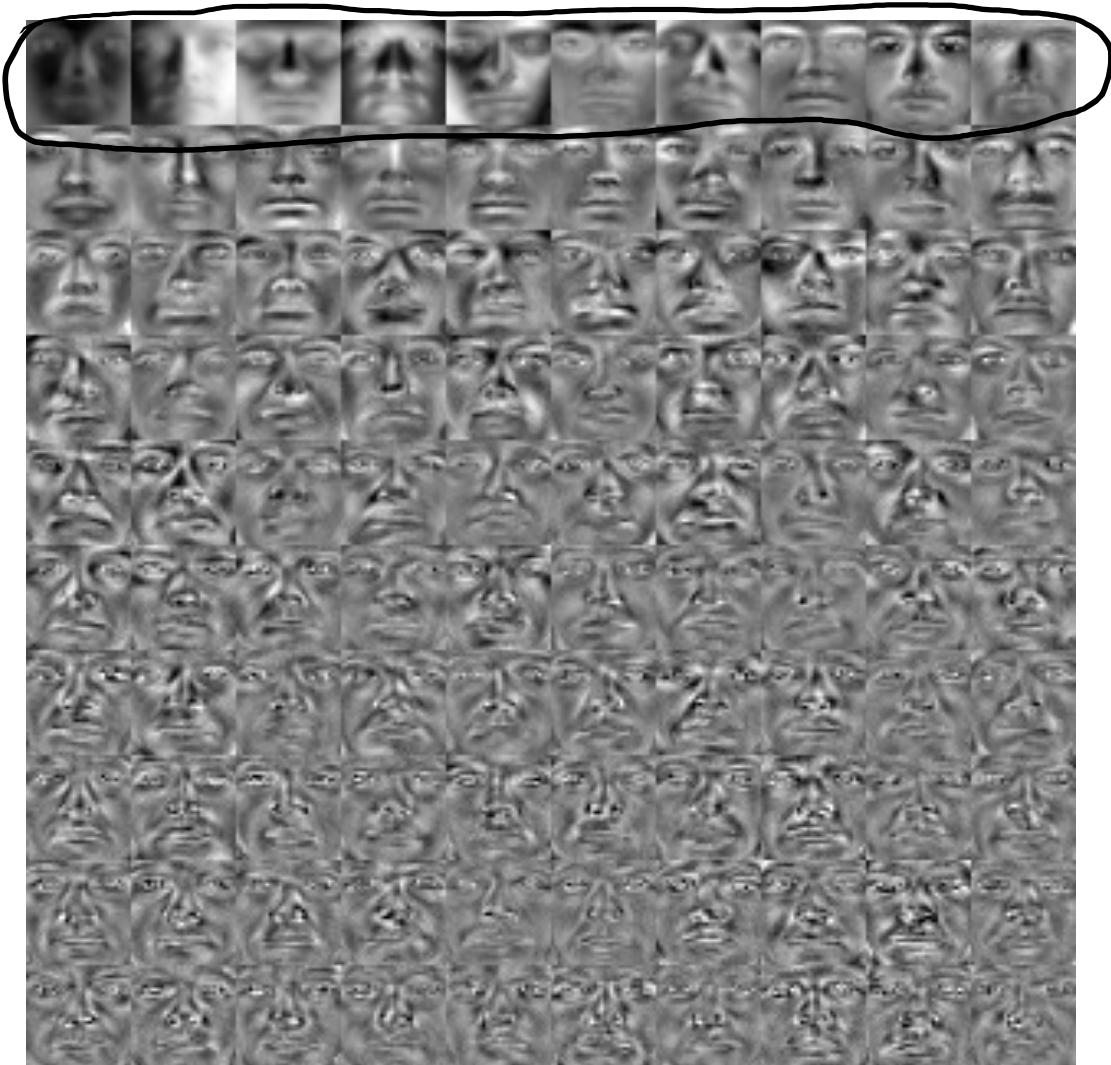


Eigenfaces

Reconstruction with $k=10$

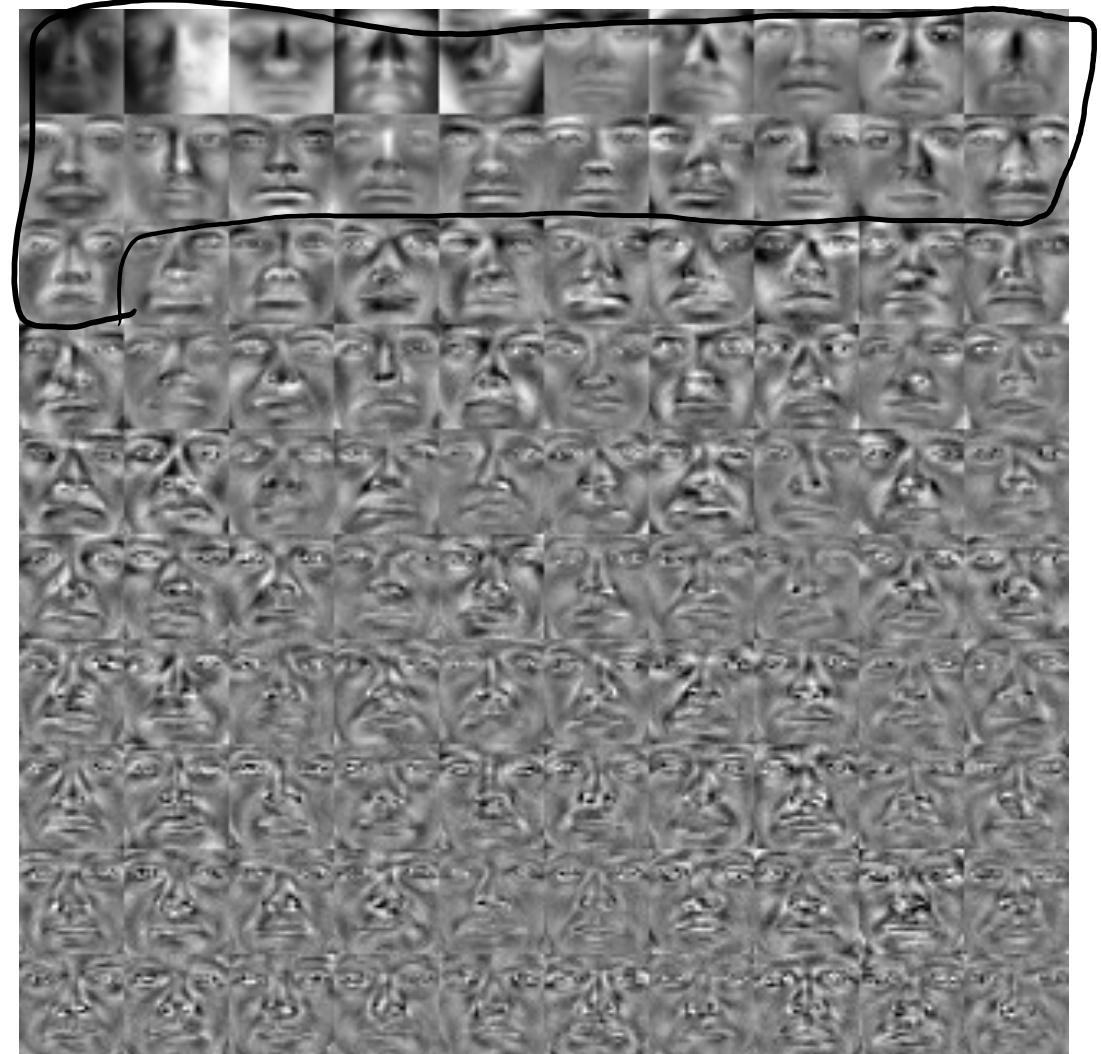


Variance explained: 85%



Eigenfaces

Reconstruction with $k=21$



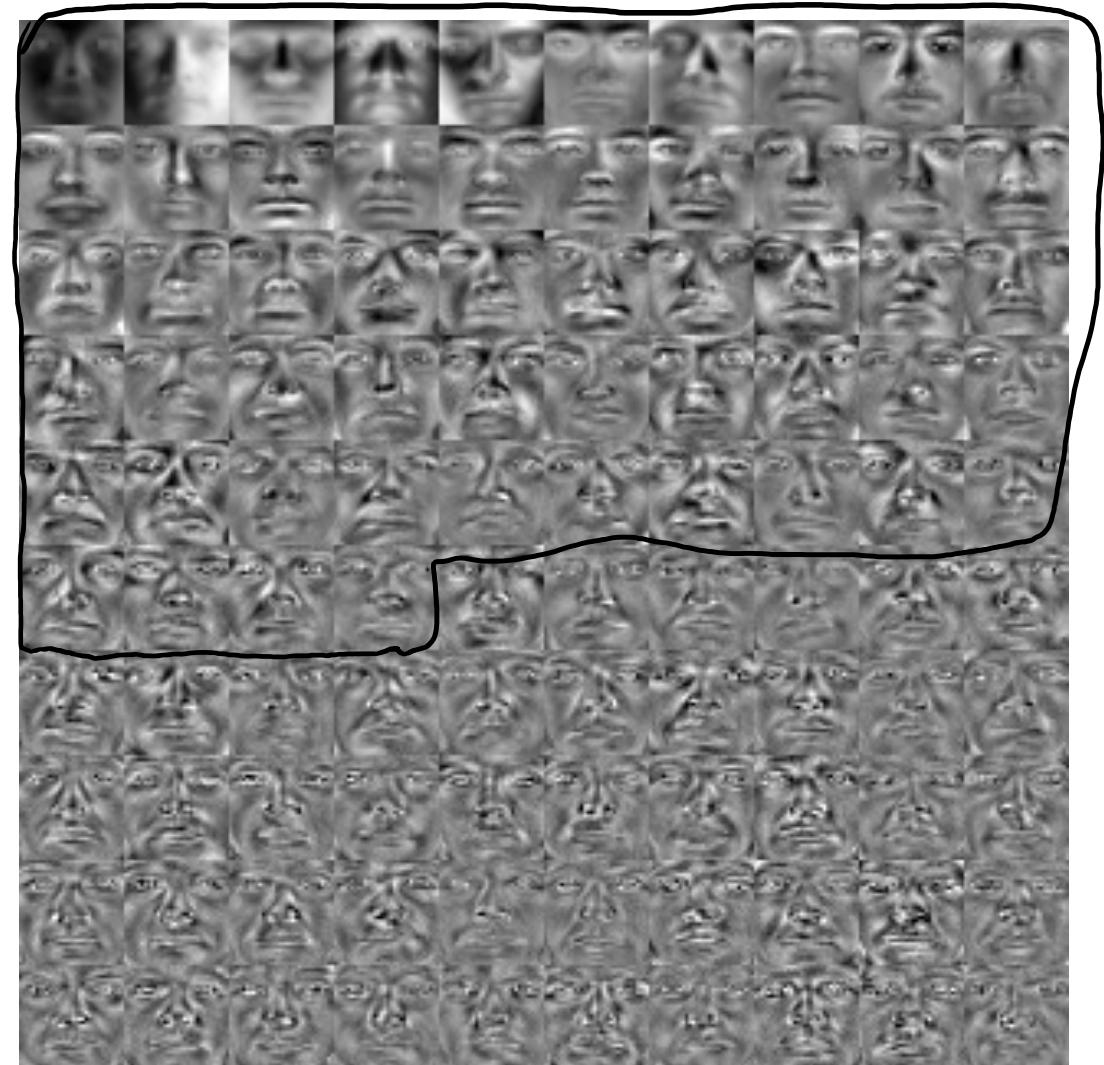
Variance explained: 90%

Eigenfaces

Reconstruction with $k=54$



Variance explained: 95%

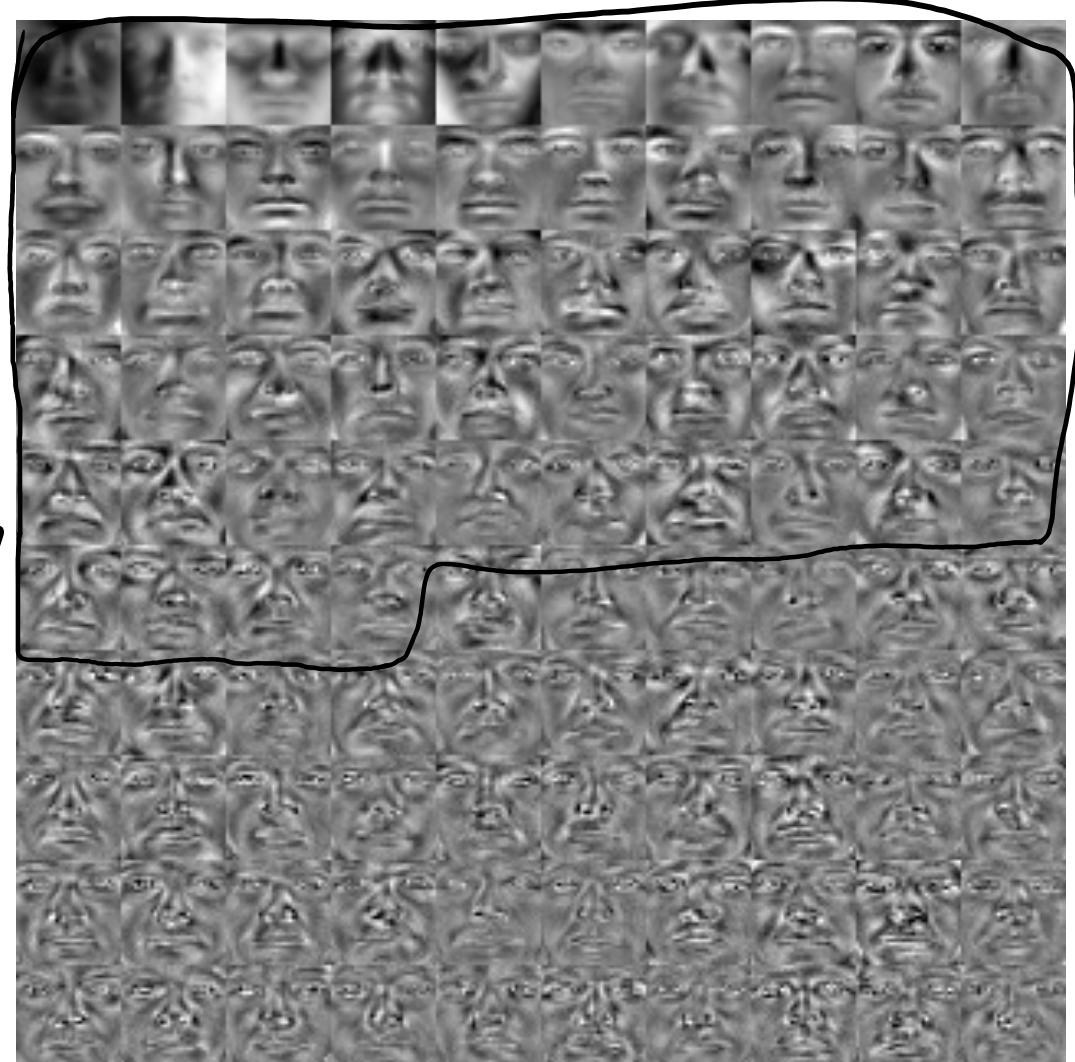


Eigenfaces

Original Images again:



Plus these
"eigenfaces"
and
the
mean,



We can replace 1024 x_i values by 54 z_i values

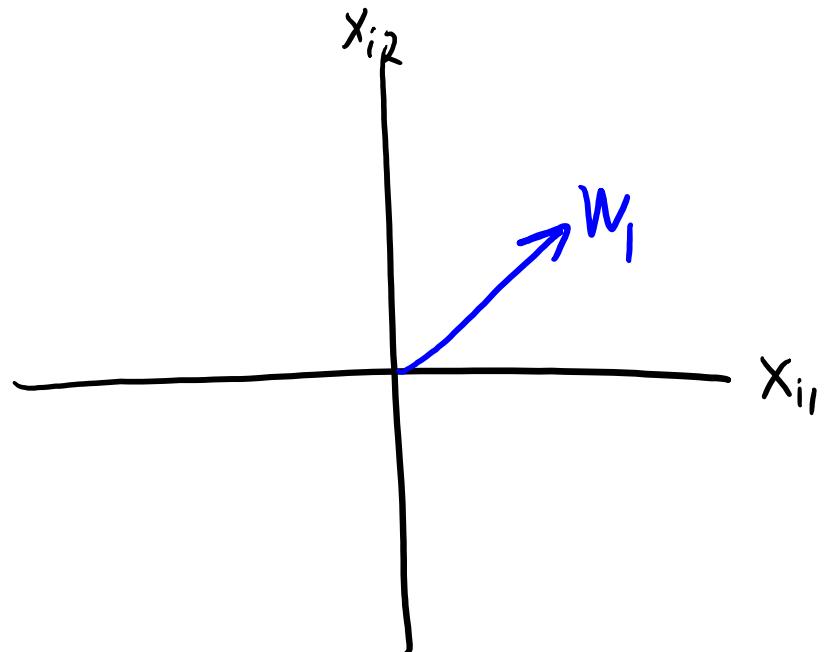
(pause)

Non-Uniqueness of PCA

- Unlike k-means, we **can efficiently find global optima** of $f(W, Z)$.
 - Algorithms coming later.
- Unfortunately, there **never exists a unique global optimum**.
 - There are actually several different sources of non-uniqueness.
- To understand these, we'll need idea of “**span**” from linear algebra.
 - This also helps explain the geometry of PCA.
 - We'll also see that **some global optima may be better than others**.

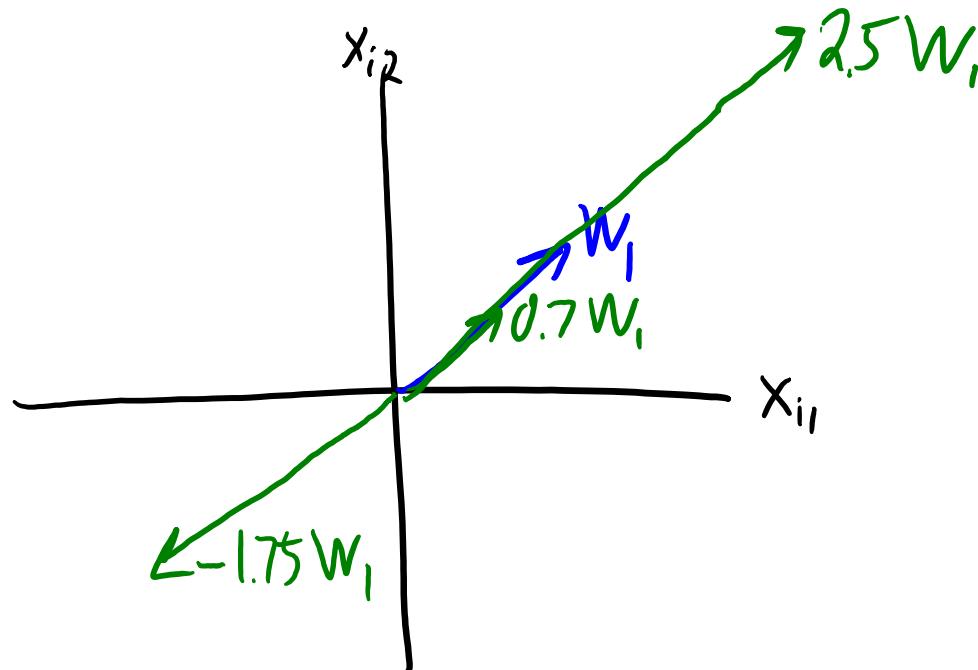
Span of 1 Vector

- Consider a single vector w_1 ($k=1$).



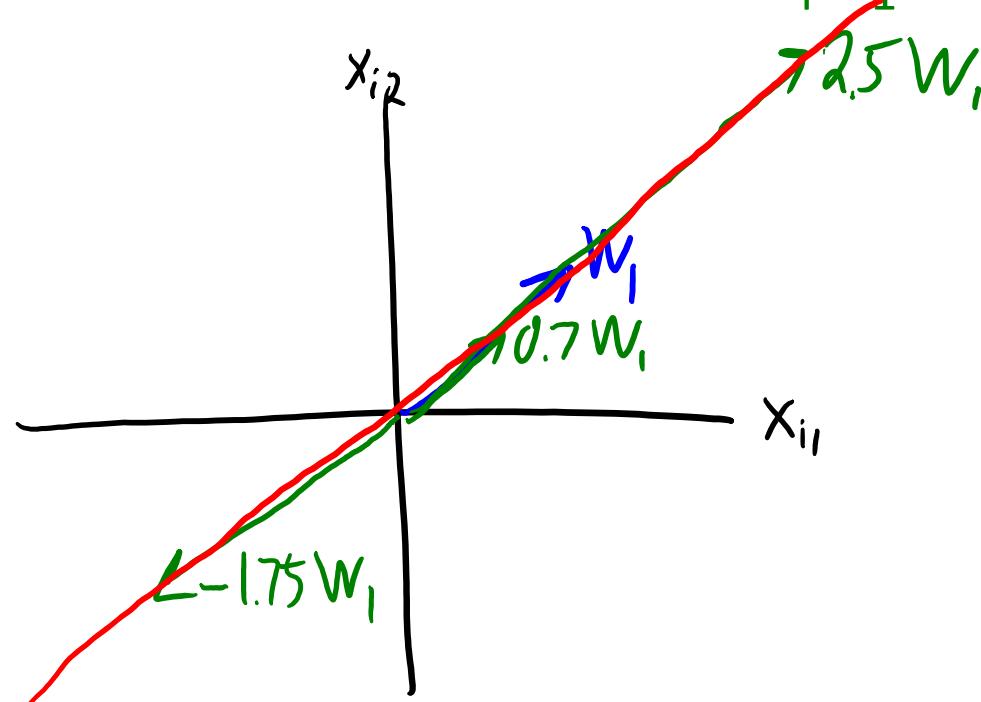
Span of 1 Vector

- Consider a single vector w_1 ($k=1$).
- The $\text{span}(w_1)$ is all vectors of the form $z_i w_1$ for a scalar z_i .



Span of 1 Vector

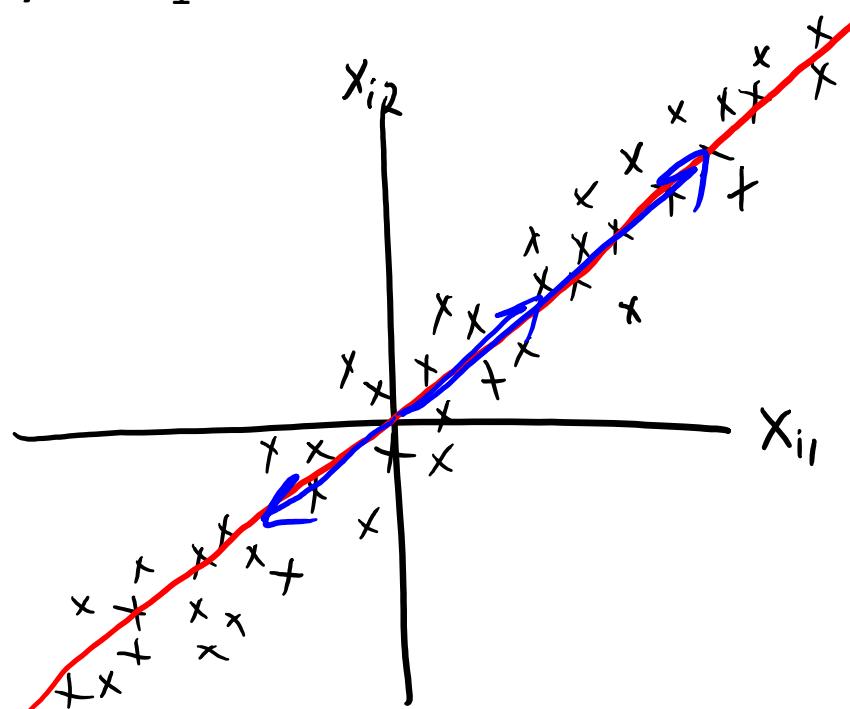
- Consider a single vector w_1 ($k=1$).
- The $\text{span}(w_1)$ is all vectors of the form $z_i w_1$ for a scalar z_i .



- If $w_1 \neq 0$, this forms a line.

Span of 1 Vector

- But note that the “span” of many different vectors gives same line.
 - Mathematically: αw_1 defines the same line as w_1 for any scalar $\alpha \neq 0$.

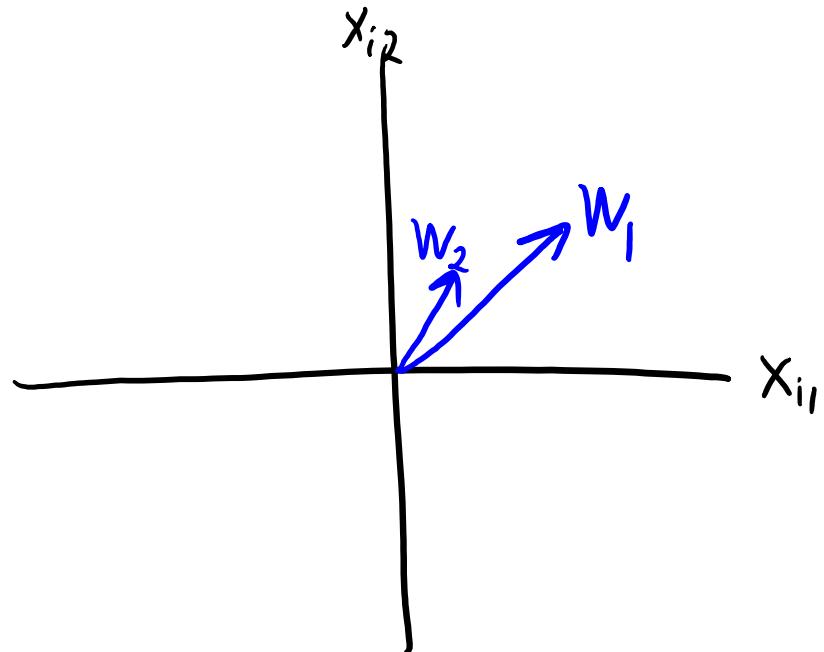


– PCA solution can only be defined up to scalar multiplication.

- If (W, Z) is a solution, then $(\alpha W, (1/\alpha)Z)$ is also a solution. $\|(\alpha W)(\frac{1}{\alpha}Z) - X\|_F^2 = \|WZ - X\|_F^2$

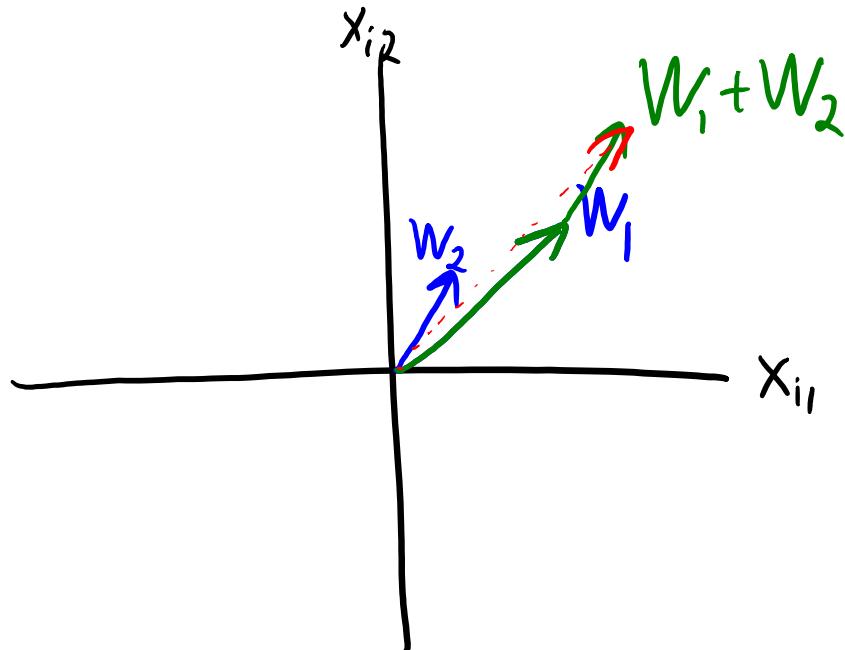
Span of 2 Vectors

- Consider two vector w_1 and w_2 ($k=2$).



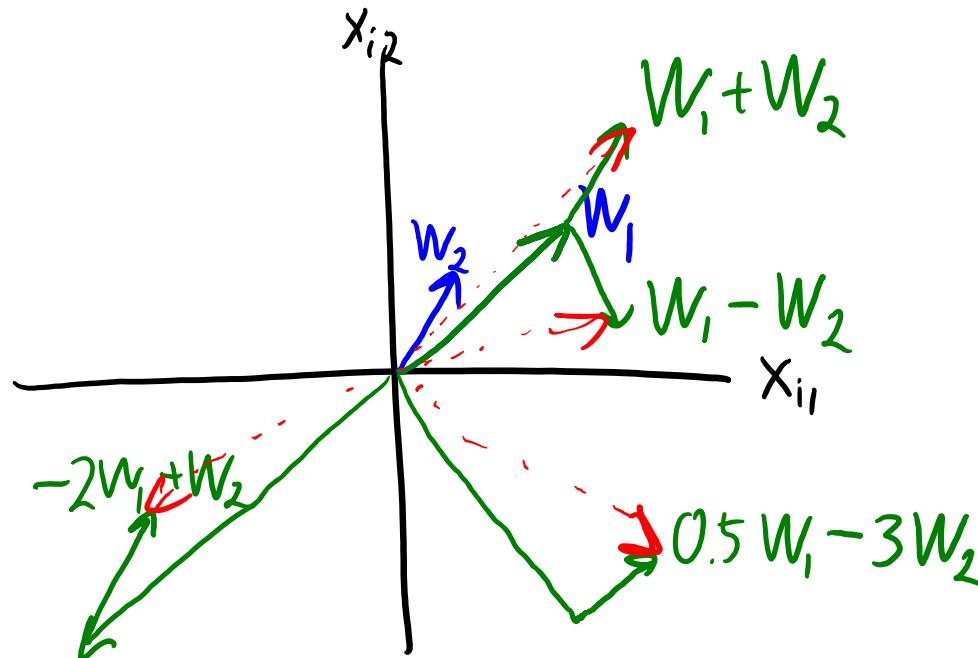
Span of 2 Vectors

- Consider two vector w_1 and w_2 ($k=2$).
 - The $\text{span}(w_1, w_2)$ is all vectors of form $z_{i1}w_1 + z_{i2}w_2$ for a scalars z_{i1} and z_{i2} .



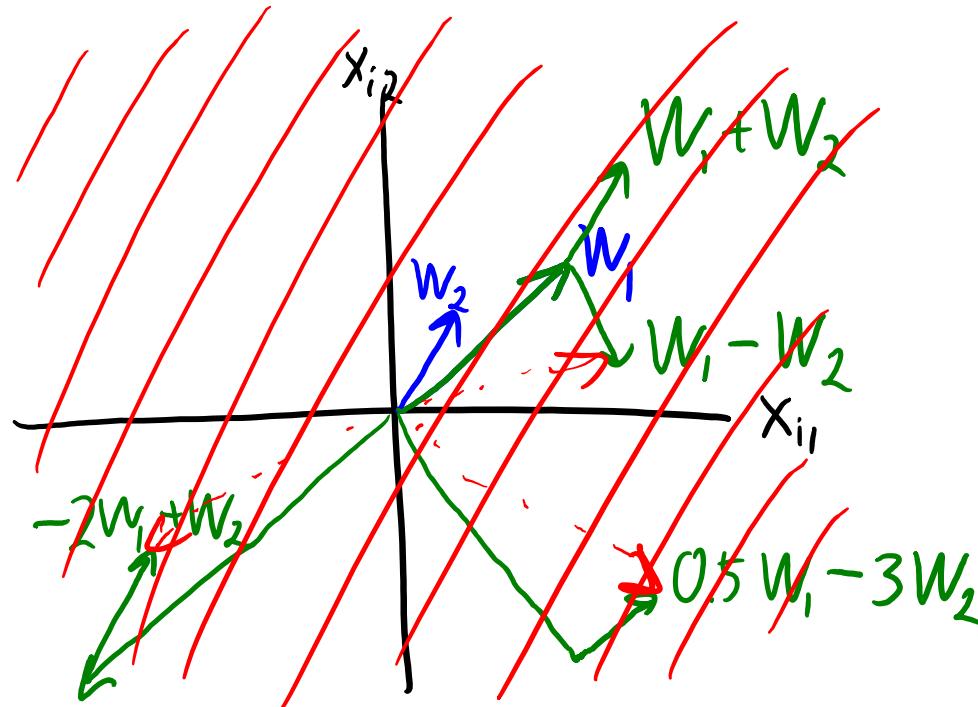
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Span of 2 Vectors

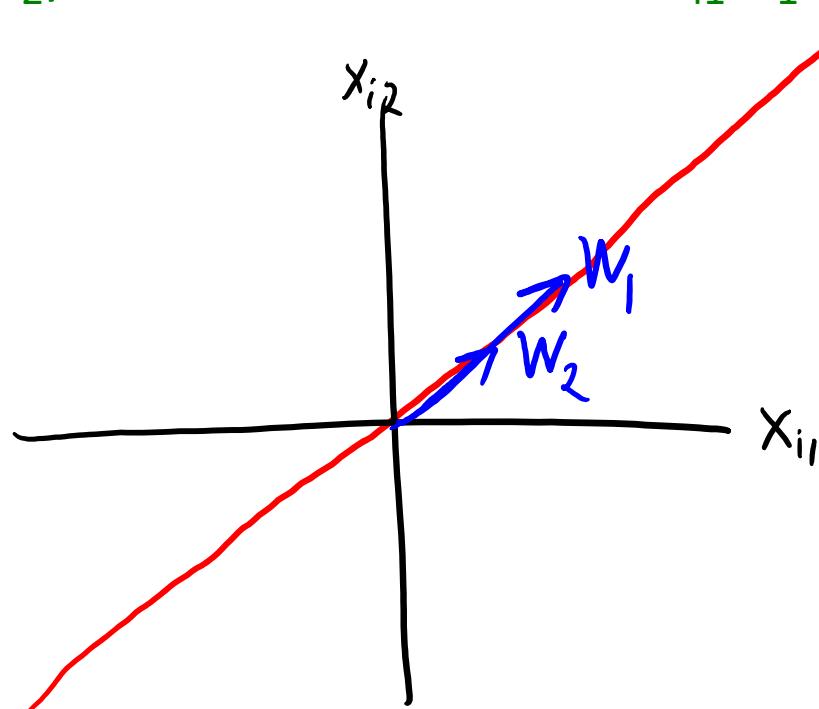
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 - The $\text{span}(w_1, w_2)$ is all vectors of form $z_{i1}w_1 + z_{i2}w_2$ for a scalars z_{i1} and z_{i2} .



- For most non-zero 2d vectors, $\text{span}(w_1, w_2)$ is a plane.
 - In the case of two vectors in R^2 , the plane will be *all* of R^2 .

Span of 2 Vectors

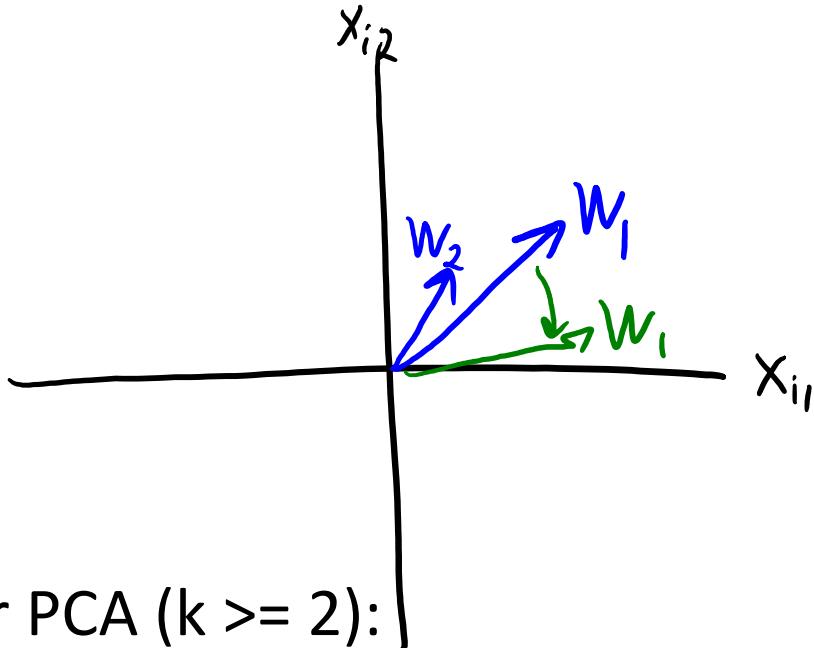
- Consider two vector w_1 and w_2 ($k=2$).
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- For most non-zero 2d vectors, $\text{span}(w_1, w_2)$ is plane.
 - Exception is if w_2 is in span of w_1 ("collinear"), then $\text{span}(w_1, w_2)$ is just a line.

Span of 2 Vectors

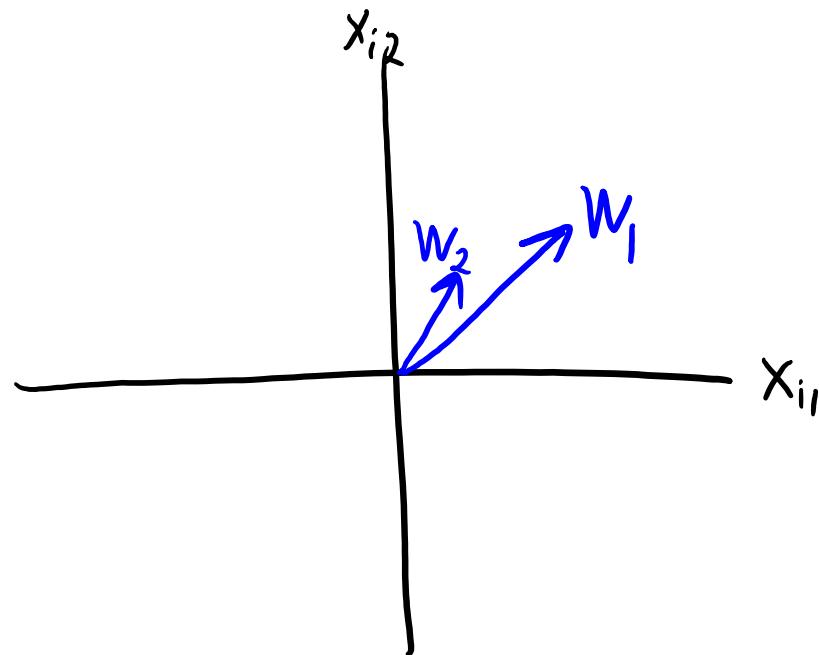
- Consider two vector w_1 and w_2 ($k=2$).
 - The $\text{span}(w_1, w_2)$ is all vectors of form $z_{i1}w_1 + z_{i2}w_2$ for a scalars z_{i1} and z_{i2} .



- New issues for PCA ($k \geq 2$):
 - We have **label switching**: $\text{span}(w_1, w_2) = \text{span}(w_2, w_1)$.
 - We can **rotate factors** within the plane (if not rotated to be collinear).

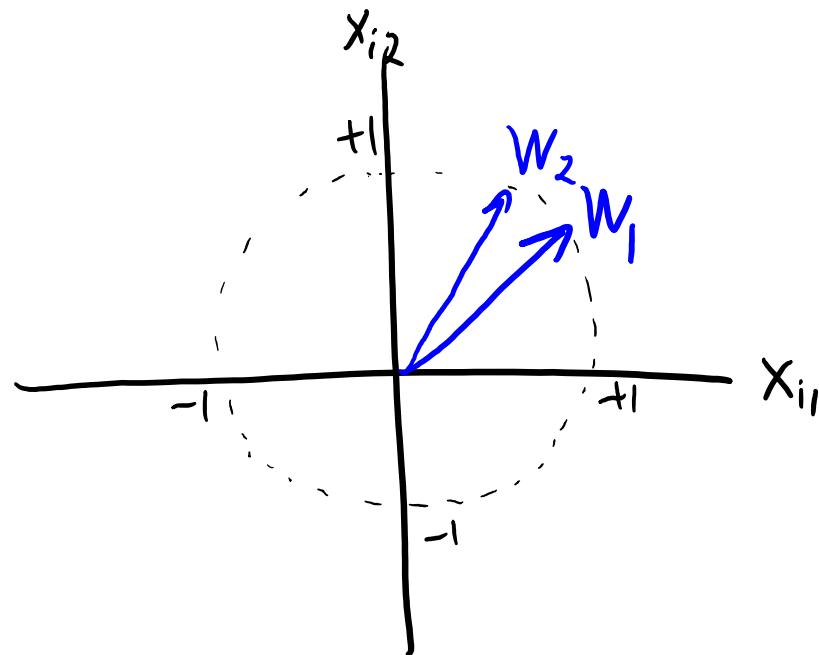
Span of 2 Vectors

- 2 tricks to make vectors defining a plane “more unique”:
 - Normalization: enforce that $\|w_1\| = 1$ and $\|w_2\| = 1$.



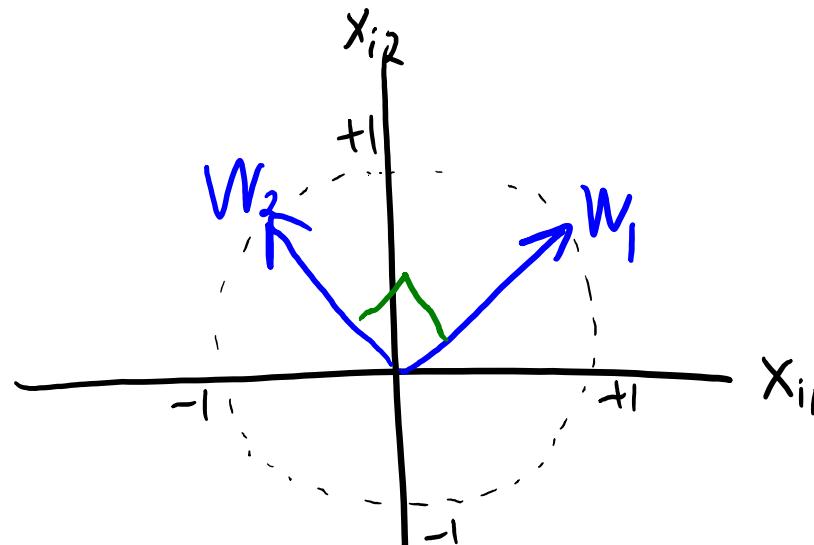
Span of 2 Vectors

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 - **Normalization**: enforce that $\|w_1\| = 1$ and $\|w_2\| = 1$.



Span of 2 Vectors

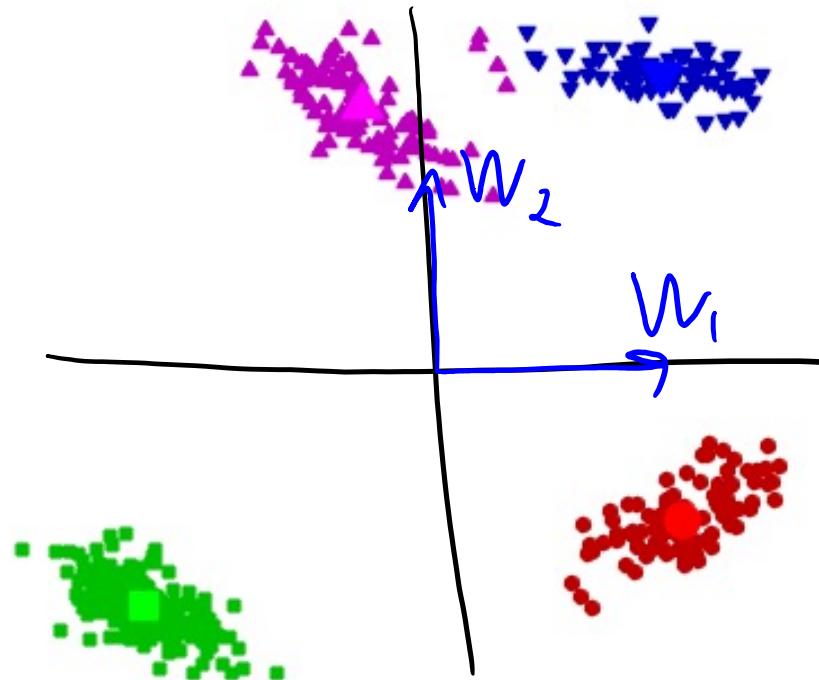
- 2 tricks to make vectors defining a plane “more unique”:
 - Normalization: enforce that $\|w_1\| = 1$ and $\|w_2\| = 1$.
 - Orthogonality: enforce that $w_1^T w_2 = 0$ (“perpendicular”).



- Now I can’t grow/shrink vectors (though I can still reflect).
- Now I can’t rotate one vector (but I can still rotate *both*).

Digression: PCA only makes sense for $k \leq d$

- Remember our clustering dataset with 4 clusters:



- It **doesn't make sense to use PCA with $k=4$** on this dataset.
 - We **only need two vectors** $[1 \ 0]$ and $[0 \ 1]$ to exactly represent all 2d points.
 - With $k=2$, I could set $Z=X$ and $W=I$ to get $X=ZW$ exactly.

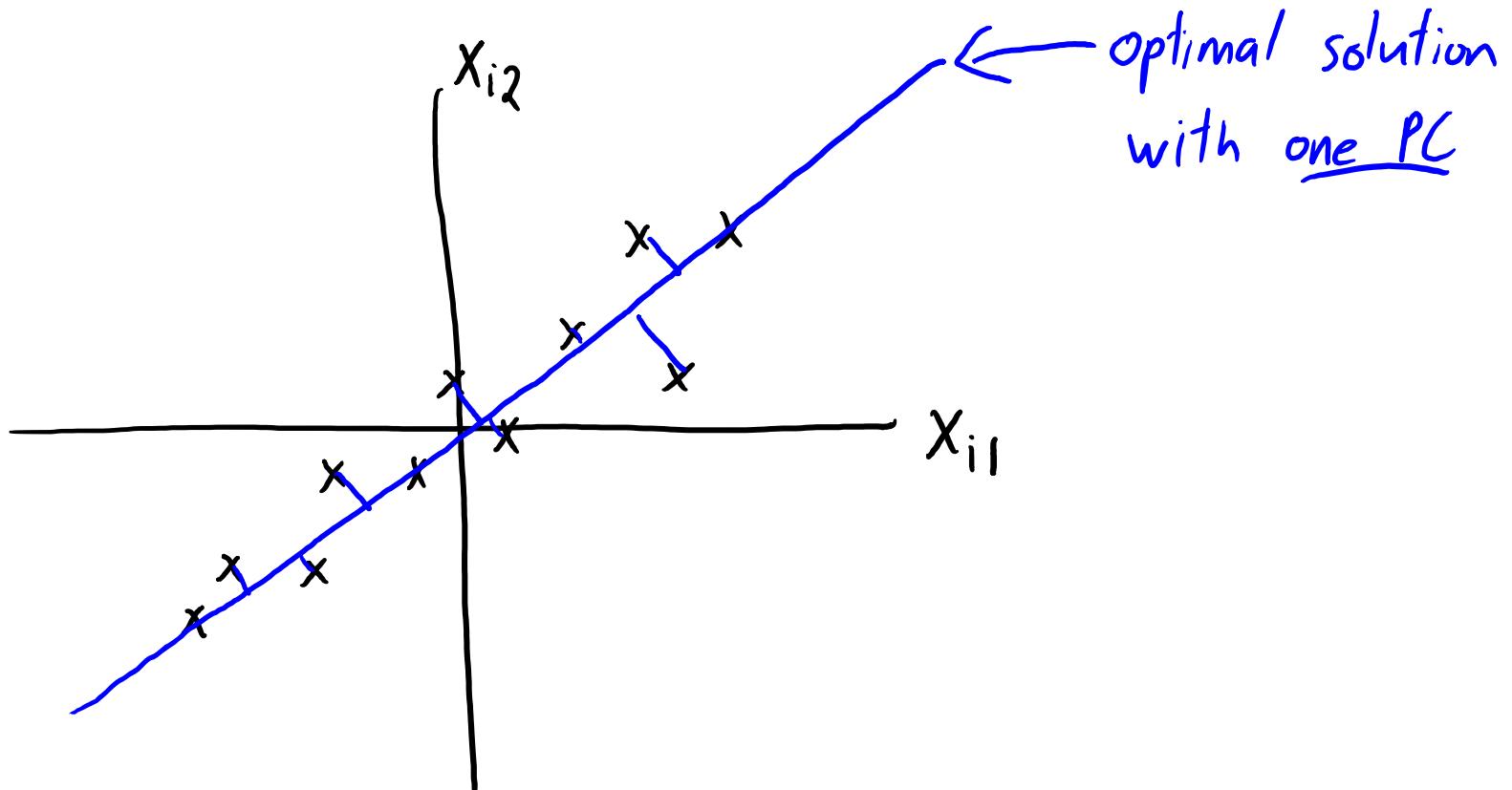
Span in Higher Dimensions

- In higher-dimensional spaces:
 - Span of 1 non-zero vector w_1 is a line.
 - Span of 2 non-zero vectors w_1 and w_2 is a plane (if not collinear).
 - Can be visualized as a 2D plot.
 - Span of 3 non-zeros vectors $\{w_1, w_2, w_3\}$ is a 3d space (if not “coplanar”).
 - ...
- This is how the W matrix in PCA defines lines, planes, spaces, etc.
 - Each time we increase ‘ k ’, we add an extra “dimension” to the “subspace”.

Making PCA Unique

- We've identified several reasons that optimal W is non-unique:
 - I can multiply any w_c by any non-zero α .
 - I can rotate any w_c almost arbitrarily within the span.
 - I can switch any w_c with any other $w_{c'}$.
- PCA implementations add constraints to make solution unique:
 - Normalization: we enforce that $\|w_c\| = 1$.
 - Orthogonality: we enforce that $w_c^T w_{c'} = 0$ for all $c \neq c'$.
 - Sequential fitting: We first fit w_1 ("first principal component") giving a line.
 - Then fit w_2 given w_1 ("second principal component") giving a plane.
 - Then we fit w_3 given w_1 and w_2 ("third principal component") giving a space.

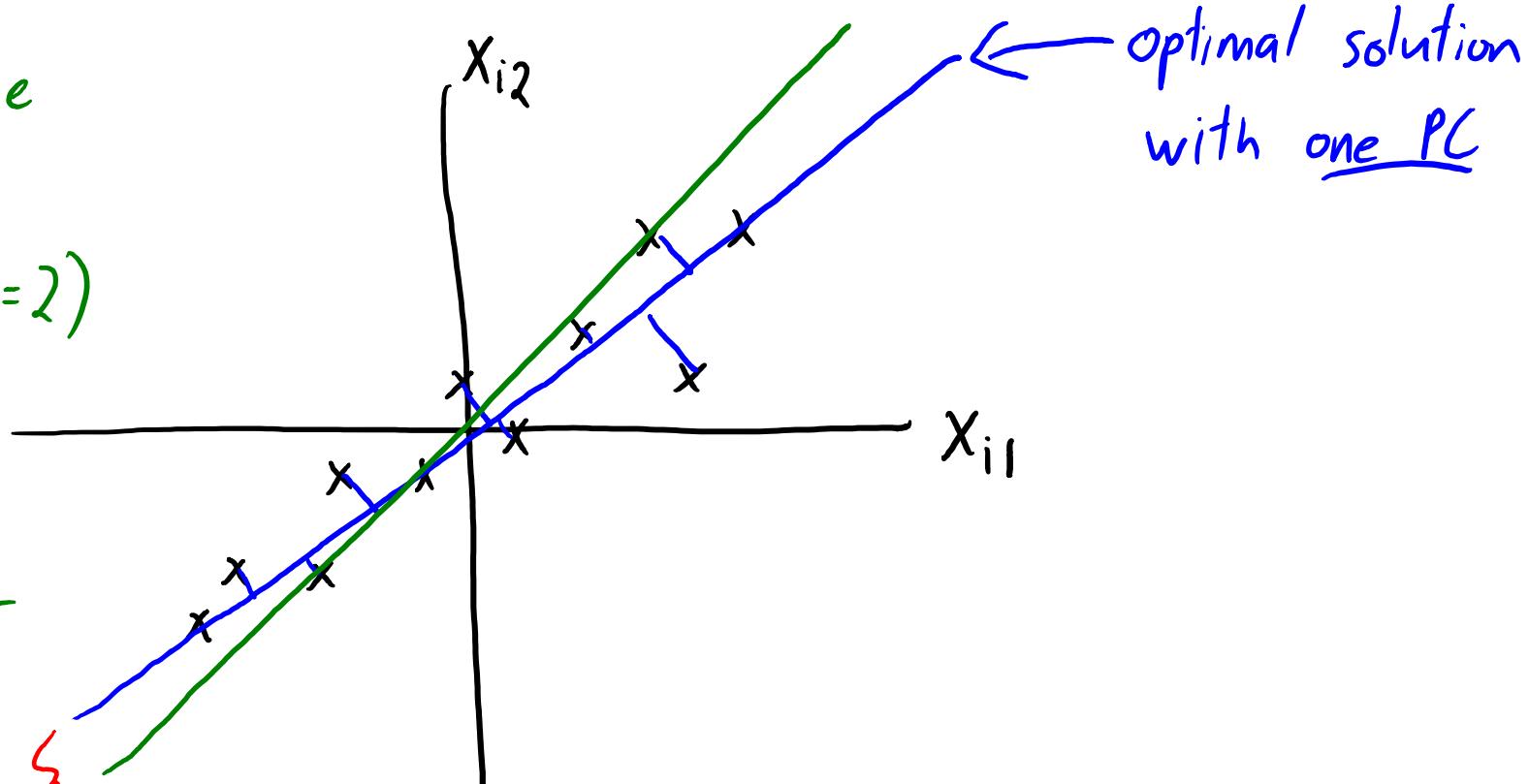
Basis, Orthogonality, Sequential Fitting



Basis, Orthogonality, Sequential Fitting

Any non-parallel line
gives optimal solution
to second PC (when $d=2$)

I can get 0 error
on every data point.



An optimal solution but not orthogonal.

(both PCs give similar information)

optimal solution
with one PC

Basis, Orthogonality, Sequential Fitting

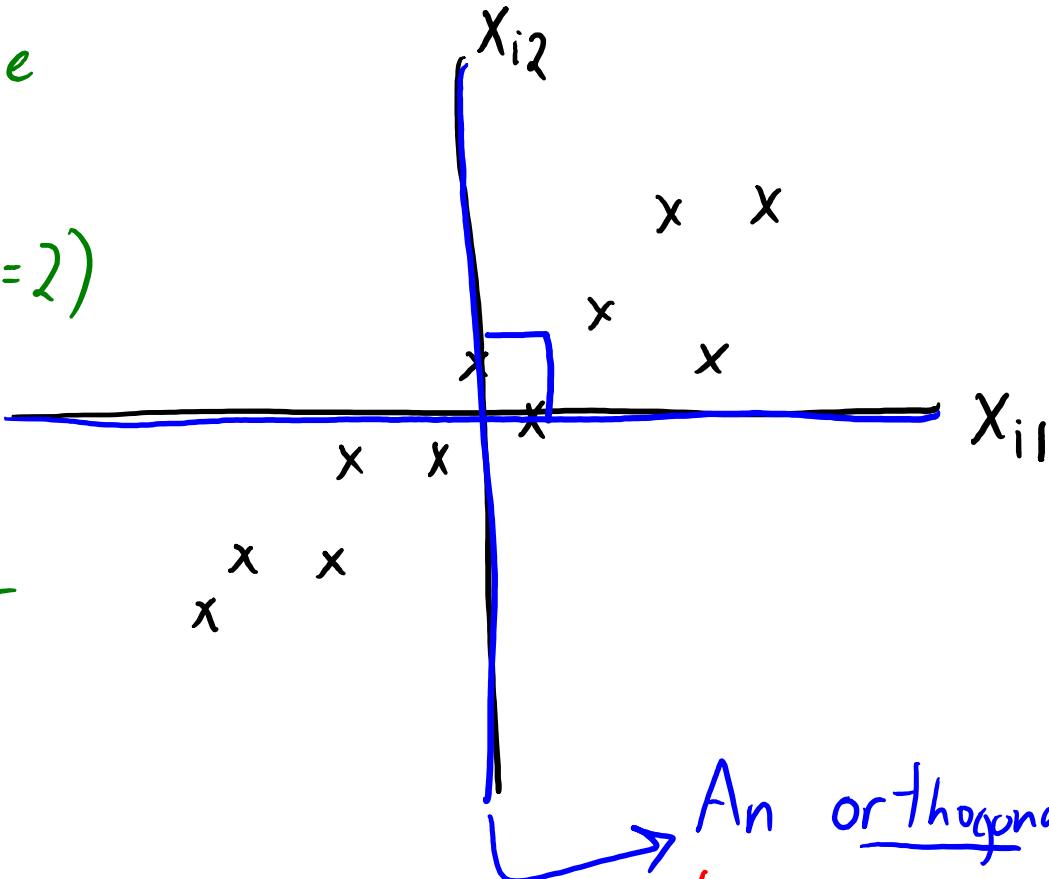
Any non-parallel line

gives optimal solution

to second PC (when $d=2$)

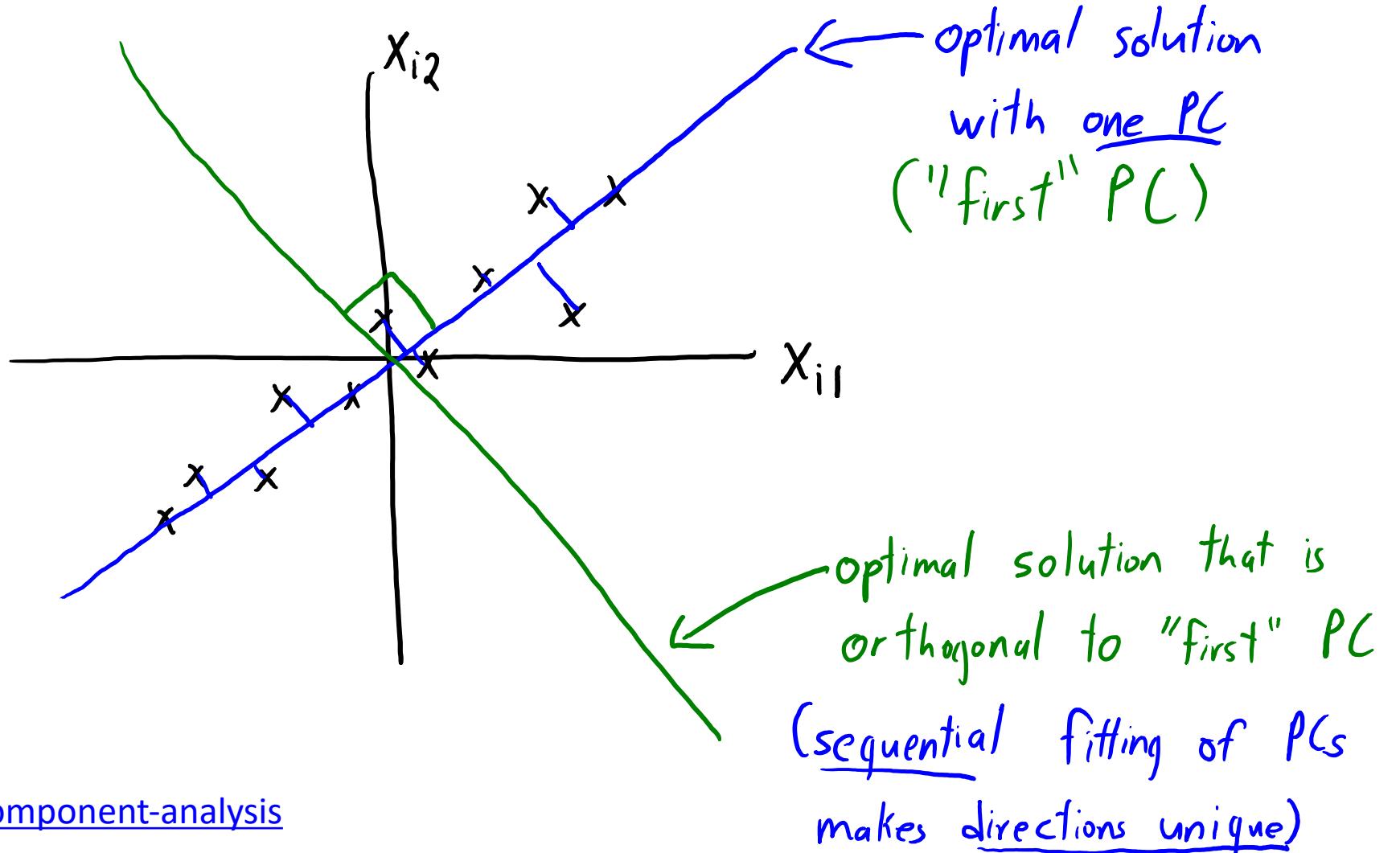


I can get 0 error
on every data point.



An orthogonal solution (PCs are not redundant)
but PCs have nothing to do with data

Basis, Orthogonality, Sequential Fitting



bonus!

PCA Computation: SVD

- How do we fit with normalization/orthogonality/sequential-fitting?
 - It can be done with the “singular value decomposition” (SVD).
 - Take CPSC 302.
- 4 lines of Python code:
 - `mu = np.mean(X, axis=0)`
 - `X -= mu`
 - `U, s, Vh = np.linalg.svd(X)`
 - `W = Vh[:k]`
- Computing Z is cheaper now:
$$Z = X W^\top (WW^\top)^{-1} = X W^\top$$
$$WW^\top = \begin{bmatrix} w_1 \\ w_2 \\ \vdots \\ w_k \end{bmatrix} \begin{bmatrix} 1 & 1 & \dots & 1 \\ w_1^\top & w_2^\top & \dots & w_k^\top \end{bmatrix}$$
$$= \begin{bmatrix} 1 & 0 & 0 & \dots & 0 \\ 0 & 1 & 0 & \dots & 0 \\ 0 & 0 & 1 & \dots & 0 \\ \vdots & \ddots & \ddots & \ddots & 0 \\ 0 & \dots & 0 & 0 & 1 \end{bmatrix} = I$$

Summary

- PCA objective:
 - Minimizes squared error between elements of X and elements of ZW .
- Choosing ‘ k ’:
 - We can choose ‘ k ’ to explain “percentage of variance” in the data.
- PCA non-uniqueness:
 - Due to scaling, rotation, and label switching.
- Orthogonal basis and sequential fitting of PCs (via SVD):
 - Leads to non-redundant PCs with unique directions.
- Next time: cancer signatures and NBA shot charts.

bonus!

Making PCA Unique

- PCA implementations add **constraints** to make solution unique:
 - **Normalization**: we enforce that $\|w_c\| = 1$.
 - **Orthogonality**: we enforce that $w_c^T w_{c'} = 0$ for all $c \neq c'$.
 - **Sequential fitting**: We first fit w_1 (“first principal component”) giving a line.
 - Then fit w_2 given w_1 (“second principal component”) giving a plane.
 - Then we fit w_3 given w_1 and w_2 (“third principal component”) giving a space.
 - ...
- Even with all this, the solution is **only unique up to sign changes**:
 - I can still replace any w_c by $-w_c$:
 - $-w_c$ is normalized, is orthogonal to the other $w_{c'}$, and spans the same space.
 - Possible fix: require that first non-zero element of each w_c is positive.
 - And this is assuming you don’t have repeated singular values.
 - In that case you can rotate the repeated ones within the same plane.