

# **STA 35C: Statistical Data Science III**

## **Lecture 23: Principal Component Analysis (cont'd)**

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# Announcement

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**Final exam** on Fri, June 6 (1:00 pm–3:00 pm) in classroom

- **Be on time:** The exam starts at 1:00 pm and ends at 3:00 pm sharp
- **Three hand-written cheat sheets allowed:** Letter-size (8.5"×11"), double-sided, brief formulas/notes
- **Calculator:** A simple (non-graphing) scientific calculator is allowed
- **No other materials** beyond the cheat sheets (no textbooks, etc.)
- **SDC accommodations:** Confirm scheduling to take on Thu, June 5 with AES ASAP

## Preparation:

- The exam is *cumulative* (Lectures 1–25)
- A practice final exam and brief answer key will be provided on the course webpage
- Office hours this week:
  - Instructor: Wed, 4:30–5:30pm ; Thu, 2:30–3:00pm
  - TA: Mon/Thu 1–2pm

# Today's topics

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## Principal component analysis (PCA)

- **Overview & intuition**
  - Objective: dimension reduction with minimal information loss
  - Intuition: projection that retains maximum variance
- **Formalism & properties**
  - Principal components (PCs)
  - PCA as a change of basis
  - Proportion of variance explained
  - Choosing number of PCs via scree plot
  - (Optional) Additional details (scaling, uniqueness, etc.)
- **Applications of PCA**

# Quick review: Unsupervised learning

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## Two branches of statistical learning:

- *Supervised learning*
  - Setup/goal: We observe  $(X, Y)$  and want to learn a function  $f : X \rightarrow Y$
  - Examples: regression, classification, ...
- *Unsupervised learning*
  - Setup/goal: We observe only  $X$  (no  $Y$ ) and aim to discover patterns or structures within  $X$
  - Examples:
    - PCA: find a few directions that capture most variation (=information) in the data
    - Clustering: identify subgroups (clusters) among observations

## Why unsupervised learning?

- We may have data only on features  $X$ ; or we want to do exploratory analysis
- Often a preliminary step before supervised tasks

# PCA: Overview & intuition

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## Problem Setup:

- We have data of  $X \in \mathbb{R}^p$ , where  $p$  is possibly large
- We want to **reduce dimension** to  $r \ll p$  while **retaining most “information”**

## PCA approach:

- Project data ( $X$ ) onto an  $r$ -dimensional subspace (spanned by  $r$  vectors)
- These  $r$  vectors (=PCs) are chosen to **capture maximum variance in  $X$**
- Unsupervised learning: no  $Y$  is used

## Outcome:

- A few linear combinations of  $X_1, \dots, X_p$  that explain most variation
- Useful for dimension reduction, model interpretation, and data visualization

## PCA illustration 1: $p = 2$ to $r = 1$

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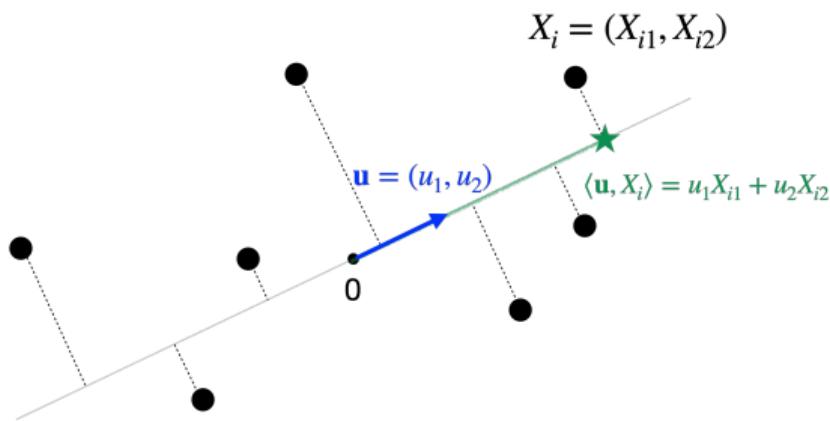


Figure: For a unit vector  $\mathbf{u} = (u_1, u_2)$ , consider the orthogonal projection of  $X_i$  onto the line spanned by  $\mathbf{u}$ , which is  $\langle \mathbf{u}, X_i \rangle = u_1 X_{i1} + u_2 X_{i2}$ . PCA finds the direction  $\mathbf{u}$  maximizing the variance of projection  $\langle \mathbf{u}, X_i \rangle$ .

### 2D $\rightarrow$ 1D projection:

- Each data point  $X_i = (x_{i1}, x_{i2})$  is mapped to  $\langle \mathbf{u}, X_i \rangle = u_1 X_{i1} + u_2 X_{i2}$
- PCA picks  $\mathbf{u}$  (with  $\|\mathbf{u}\| = 1$ ) that *maximizes* the variance of  $\langle \mathbf{u}, X_i \rangle$ ,  $\frac{1}{n} \sum_{i=1}^n \langle \mathbf{u}, X_i \rangle^2$
- Geometrically, the "major axis" of the data cloud is identified

## PCA example: $p = 2$ data to $r = 1$

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### Example

Consider a small 2D dataset:

$$\mathcal{X} = \{(-2, -1), (0, 0), (2, 1)\}.$$

These three points lie on the line spanned by  $(2, 1)$ .

**Projection:** For a unit vector  $\mathbf{u} = (u_1, u_2)$ , the projection of  $X_i = (x_{i1}, x_{i2})$  onto (the line spanned by)  $\mathbf{u}$  is

$$\langle \mathbf{u}, X_i \rangle = u_1 x_{i1} + u_2 x_{i2}.$$

**Key idea:** PCA finds  $\mathbf{u}$  that maximizes the variance of these projected values  $\langle \mathbf{u}, X_i \rangle$ .

Observe that

- If  $\mathbf{u} = (1, 0)$ , the variance in this direction is  $\frac{1}{3}((-2)^2 + 0^2 + 2^2) = \frac{8}{3}$ .
- If  $\mathbf{u} = (0, 1)$ , the variance in this direction is  $\frac{1}{3}((-1)^2 + 0^2 + 1^2) = \frac{2}{3}$ .
- If  $\mathbf{u} = \frac{1}{\sqrt{5}}(2, 1)$ , the variance in this direction is  $\frac{1}{3}((-2\sqrt{5})^2 + 0^2 + (\sqrt{5})^2) = \frac{10}{3}$  (the maximum).

Hence  $\mathbf{u}^* = \frac{1}{\sqrt{5}}(2, 1)$  is the PCA direction.

## PCA illustration 2: $p = 3$ to $r = 2$

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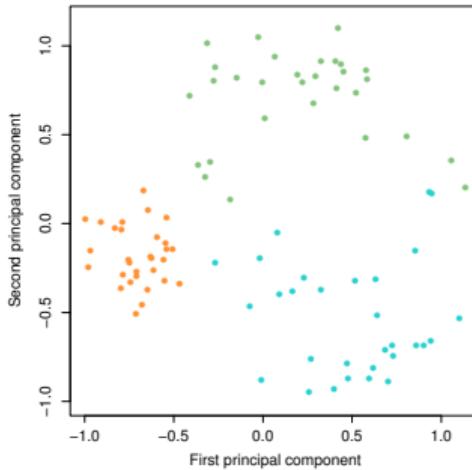
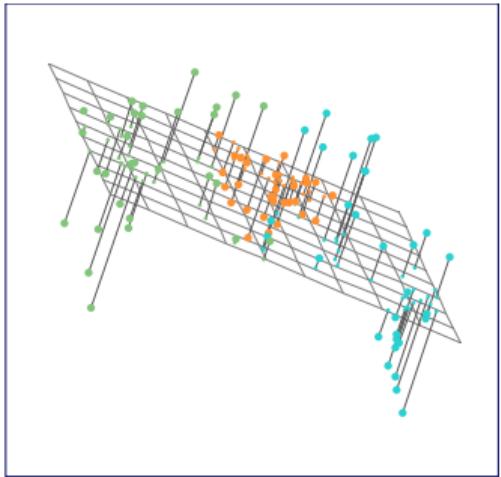


Figure: Ninety observations in  $\mathbb{R}^3$ . **Left:** The first two PC directions span a plane that best fits the data, minimizing total squared distance. **Right:** Data are “flattened” onto that 2D plane, forming their PC scores [JWHT21, Figure 12.2].

### Key idea in higher dimension:

- Find a subspace of dimension  $r$  that capture maximal variance (=minimizing residuals)
- $\mathbf{u}_1$  is the top PC direction,  $\mathbf{u}_2$  is second, etc., each orthogonal

## PCA: Formulation ( $p = 2$ , $r = 1$ )

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**Assumption:** Data is *centered*, i.e.

$$\mathbb{E}[X] = \frac{1}{n} \sum_{i=1}^n X_i = 0$$

**First principal component direction** = the direction  $\mathbf{u} = (u_1, u_2)$  that solves

$$\text{maximize } \frac{1}{n} \sum_{i=1}^n (\mathbf{u} \cdot \mathbf{x}_i)^2 \quad \text{subject to} \quad \|\mathbf{u}\| = \sqrt{u_1^2 + u_2^2} = 1$$

- The *variance* of  $X$  along  $\mathbf{u}$  is

$$\text{Var}(\langle \mathbf{u}, X \rangle) = \mathbb{E} \left[ (\langle \mathbf{u}, \mathbf{X} \rangle - \underbrace{\mathbb{E} \langle \mathbf{u}, \mathbf{X} \rangle}_{=0})^2 \right] = \frac{1}{n} \sum_{i=1}^n (\mathbf{u} \cdot \mathbf{x}_i)^2 = \frac{1}{n} \sum_{i=1}^n (u_1 x_{i1} + u_2 x_{i2})^2.$$

- Geometrically, the solution is the “major axis” in  $\mathbb{R}^2$  that explains the largest spread

# PCA: General formulation ( $p \geq 2, r \geq 1$ )

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**First PC:** a unit vector  $\mathbf{u}_1 \in \mathbb{R}^p$  that maximizes variance, i.e.,

$$\mathbf{u}_1 = \operatorname{argmax}_{\|\mathbf{u}\|=1} \frac{1}{n} \sum_{i=1}^n (\mathbf{u} \cdot \mathbf{x}_i)^2$$

**Second PC:** a unit vector  $\mathbf{u}_2 \in \mathbb{R}^p$  maximizing variance, subject to being orthogonal to  $\mathbf{u}_1$ ,

$$\mathbf{u}_2 = \operatorname{argmax}_{\substack{\|\mathbf{v}\|=1 \\ \langle \mathbf{v}, \mathbf{u}_1 \rangle = 0}} \frac{1}{n} \sum_{i=1}^n (\mathbf{v} \cdot \mathbf{x}_i)^2$$

- $\mathbf{u}_1 \perp \mathbf{u}_2 \implies$  the random variables  $Z_1 = \langle \mathbf{u}_1, X \rangle$  and  $Z_2 = \langle \mathbf{u}_2, X \rangle$  are *uncorrelated*
- Subsequent PCs  $\mathbf{u}_3, \dots, \mathbf{u}_p$  are defined analogously, each orthogonal to all preceding PCs

**Interpretation:**

- The  $k$ -th PC is orthogonal to all prior ones, ensuring uncorrelatedness among the PC scores
- (Optional) The PC directions correspond to the eigenvectors of the sample covariance matrix

## PCs as linear combinations (change of basis)

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If  $\mathbf{u}_1, \dots, \mathbf{u}_p$  are PC directions, the  $k$ -th **PC score** for observation  $i$  is

$$Z_{ik} = \langle \mathbf{u}_k, X_i \rangle = \sum_{j=1}^p u_{k,j} X_{i,j}.$$

- We can write  $X_i$  as a combination of the  $\mathbf{u}_k$  basis:

$$X_i = \sum_{j=1}^p X_{ij} \mathbf{e}_j = \sum_{k=1}^p Z_{ik} \mathbf{u}_k$$

- For dimension reduction, we might keep only  $Z_{i1}, \dots, Z_{ir}$  for  $r \ll p$ , compressing the data

(Optional):

- $(Z_1, \dots, Z_p)$  is just a linear transformation of  $(X_1, \dots, X_p)$
- In matrix form,  $Z = X U$ , where  $U$  is the orthonormal matrix whose columns are  $\mathbf{u}_1, \dots, \mathbf{u}_p$

## PCs as linear combinations (change of basis)

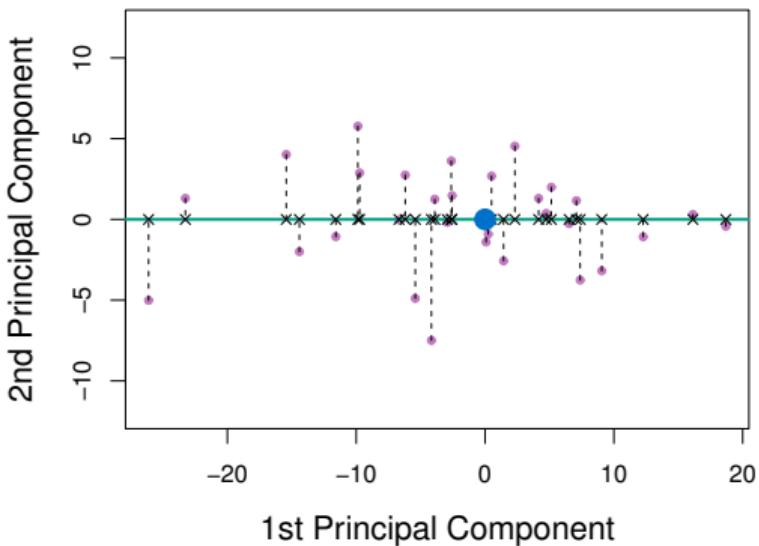
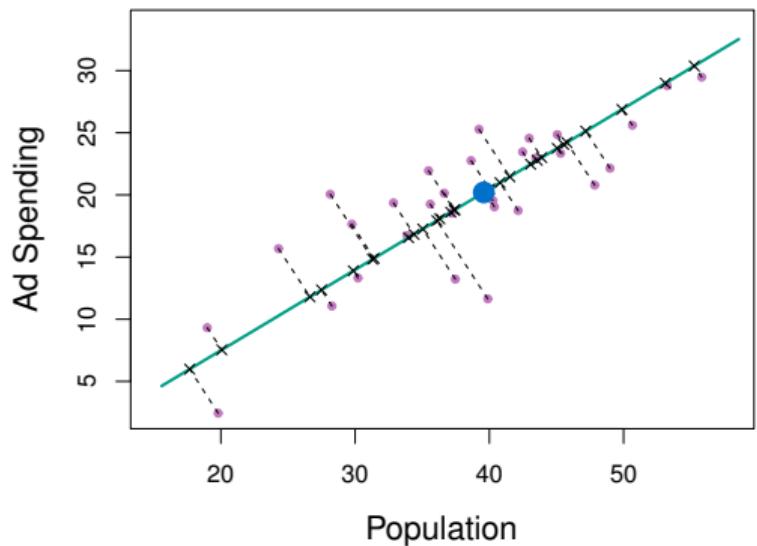


Figure: A subset of the `advertising` data, with the mean `pop` and `ad` budgets shown as a blue circle. **Left:** The first principal component direction (green) captures the greatest data variation and defines the line that best fits all observations (distances shown by dashed segments). **Right:** The plot is rotated so that this principal component aligns with the horizontal x-axis. [JWHT21, Figure 6.15].

## Pop-up quiz #1: Basic PCA understanding

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**Question:** Which statement about PCA is **false**?

- A) PCA is unsupervised, using only  $\{X_j\}$ , not  $Y$ .
- B) The first principal component is the direction in predictor space along which the projected data has the largest variance.
- C) The second principal component must be found by maximizing projected variance with no extra constraint.
- D) PCA can serve as dimension reduction by keeping only a few top PCs capturing most variance.

**Answer:** (C) is false.

- For the second principal component, there is an extra requirement that it is *uncorrelated* (and hence orthogonal, in geometric terms) to the first principal component.

## Pop-up quiz #2: Interpreting the first principal component

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**Question:** Suppose you have  $p$  predictors and you compute the first principal component. Which choice **best describes** how to interpret that first component?

- A) It is always the average of all the predictors, so it has little to do with variance.
- B) It is the unit-length direction that maximizes how spread out (variable) the data is after projecting onto that direction.
- C) It represents a decision boundary for separating classes in your dataset.
- D) It is guaranteed to pass exactly through every data point if we use all observations.

**Answer:** (B) is correct.

- The first principal component is the direction along which the data points show the greatest variance; it does not necessarily pass through every data point, nor does it reflect a classification decision boundary.

# Proportion of variance explained (PVE)

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**Question:** If we only keep  $r$  PCs, how much total variance remains?

- For centered data, *total variance* is

$$\text{Var}(X) = \frac{1}{n} \sum_{i=1}^n \|X_i\|^2 = \frac{1}{n} \sum_{i=1}^n \sum_{j=1}^p x_{ij}^2 = \sum_{j=1}^p \text{Var}(X_j)$$

- The variance explained by the  $k$ -th PC is

$$\text{Var}(\langle \mathbf{u}_k, X \rangle) = \frac{1}{n} \sum_{i=1}^n z_{ik}^2 \quad \text{where} \quad z_{ik} = \langle \mathbf{u}_k, X_i \rangle$$

- The **proportion of variance explained** (PVE) by the  $k$ -th PC is

$$\text{PVE}_k = \frac{\sum_{i=1}^n z_{ik}^2}{\sum_{i=1}^n \sum_{j=1}^p x_{ij}^2} = \frac{\sum_{i=1}^n \|z_{ik}\mathbf{u}_k\|^2}{\sum_{i=1}^n \|X_i\|^2}$$

- The **cumulative PVE** for the first  $r$  PCs is

$$\text{PVE}_{1:r} = \sum_{k=1}^r \text{PVE}_k = 1 - \frac{\sum_{i=1}^n \left\| X_i - \sum_{k=1}^r z_{ik} \mathbf{u}_k \right\|^2}{\sum_{i=1}^n \|X_i\|^2} = 1 - \frac{\text{RSS}}{\text{TSS}}$$

# Scree plot: PVE vs. number of PCs

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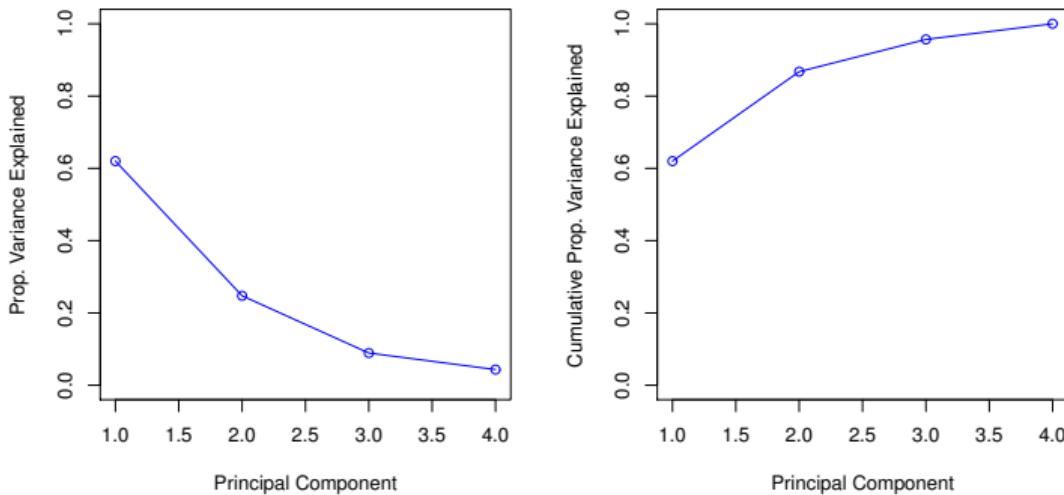


Figure: A scree plot for the [USArrests](#) data. **Left:** proportion of variance explained by each PC. **Right:** cumulative PVE [JWHT21, Figure 12.3].

## Scree plot:

- Plot PVE or cumulative PVE vs. PC index  $k$
- Often look for an “elbow” beyond which additional PCs yield minimal gains

# Scree plot: How many PCs to retain?

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## Trade-off:

- Smaller dimension  $r$  is easier to interpret and visualize
- Larger  $r$  retains more variance in the data

## Question: How many principal components do we need?

- No universal formula for the “best”  $r$
- Typically choose  $r$  so the **cumulative PVE** is “high enough,” or identify an “elbow” in the scree plot
- Larger  $r$  retains more variance (less information loss) but can be less interpretable

**In practice, use scree plot** to find an “elbow” and retain the PCs on the left

# Choosing the number of PCs using scree plot example

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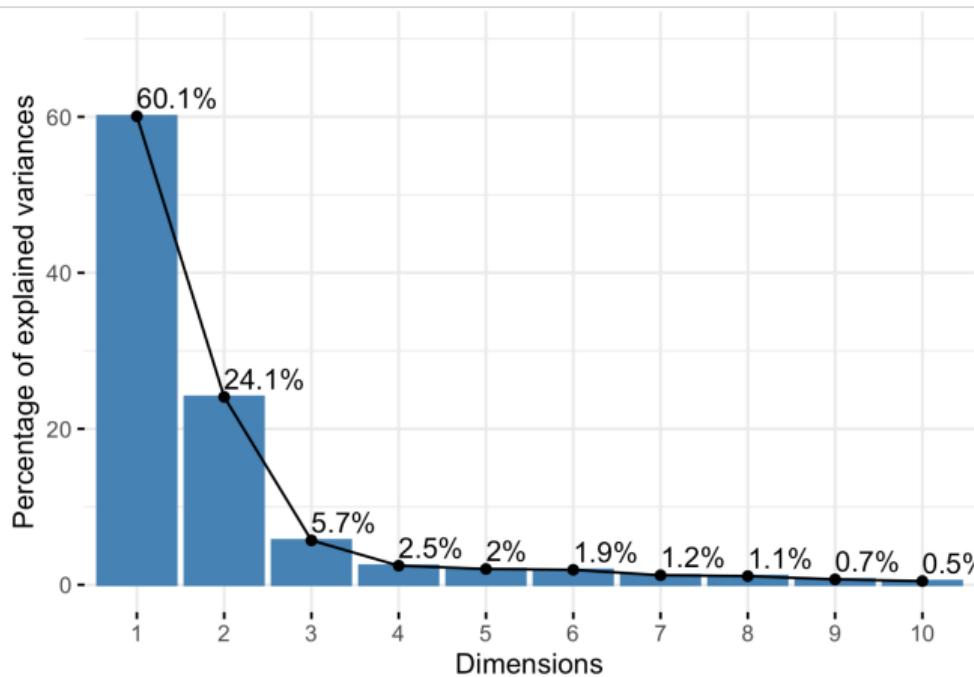


Figure: A scree plot from `mtcars` dataset in R. The elbow appears to occur at the third principal component, which suggests keeping the first three components (source: [Statistics Globe](#)).

## (Optional) Additional PCA details

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### Scaling variables?

- If predictors have very different scales (e.g. height in cm vs. income in \$), standardizing them to unit variance can drastically alter PCA directions
- Whether to scale depends on context: if raw scales matter, do not standardize; if you want each feature to contribute equally, do scale

### Uniqueness:

- Principal component directions are unique up to a sign ( $\mathbf{u}$  vs.  $-\mathbf{u}$ )
- This sign usually does not affect interpretation, so software packages pick a sign convention automatically

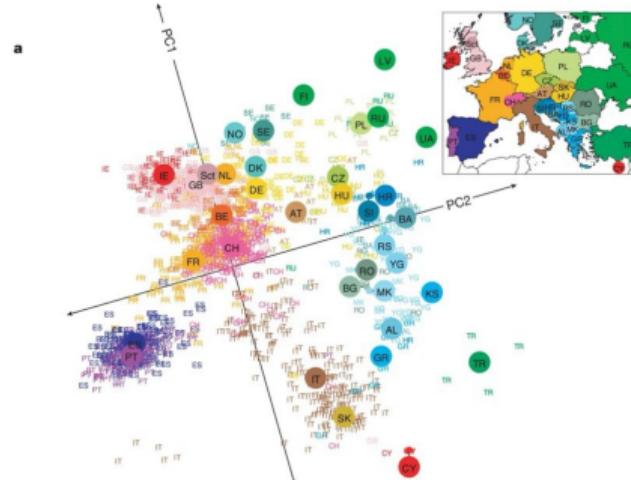
### Computation:

- Solve for eigenvectors/eigenvalues of the sample covariance (or correlation) matrix
- In R: `prcomp(..., scale=TRUE)` or `princomp(...)`

# PCA application in high-dimensional genomics

**Example:** Genomics data [NJB<sup>+</sup>08]

- 1,387 individuals from Europe, each with genotype data at 197,146 loci
- Apply PCA → reduce dimension from  $p = 197k$  to 2 principal components
- Two PCs remarkably recapitulate Europe's geography in "genetic space," demonstrating how PCA can drastically compress data while still capturing meaningful structure



**Figure:** First two principal components of genetic variation among 1,387 Europeans. Small colored points are individuals; large dots mark country medians in PC1–PC2 space [NJB<sup>+</sup>08, Figure 1-a].

# PCA application in image compression

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**Example:** Compressing a grayscale image via PCA

- Original image has  $372 \times 492$  pixels, each a grayscale intensity in  $[0, 255]$
- The image is partitioned into  $12 \times 12$  blocks, so each block is a  $12 \times 12 = 144$ -dimensional “vector”
- There are  $N = \frac{372}{12} \times \frac{492}{12} = 1271$  such vectors (observations)
- Apply PCA with rank  $r \in \{1, 3, 6, 16, 60\}$  for the dimension reduction

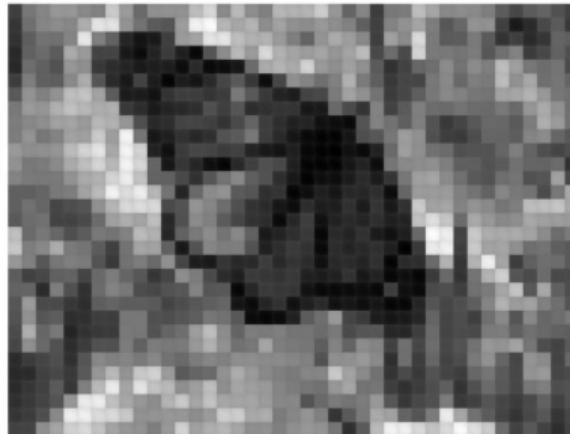
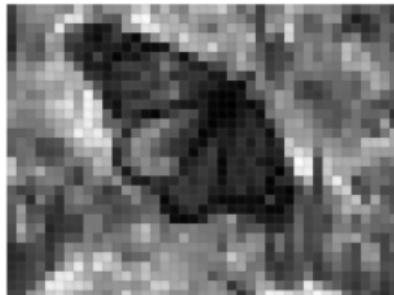


Figure: Compressing an image by PCA. **Left:** original image. **Right:** PCA rank-1 approximation. With  $r = 1$ , almost all details are lost, but the main global contrast is still visible.

## PCA application in image compression (cont'd)



Original



Rank-1 PCA ( $r = 1$ )



Rank-3 PCA ( $r = 3$ )



Rank-6 PCA ( $r = 6$ )



Rank-16 PCA ( $r = 16$ )



Rank-60 PCA ( $r = 60$ )

Figure: PCA-based image compression. Larger  $r$  yields better reconstruction quality.

# Wrap-up: Takeaways

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## Principal Component Analysis (PCA):

- Finds a few PC directions that capture maximum variance in the data
- The first few PCs often capture most of the total variation, enabling dimension reduction
- PCA is *unsupervised*, commonly used for exploratory analysis or as a pre-processing step

## Proportion of Variance Explained (PVE):

- Quantifies how much of the total variance is retained by a chosen number  $r$  of PCs
- A scree plot of PVE vs. PC index can guide how many PCs to keep

## Additional remarks:

- In R, use `prcomp(...)` or `princomp(...)`
- Predictor scaling can affect PCA
- Once you learn linear algebra & eigendecomposition, the definitions and details of PCA will become much clearer

# References

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