

# Unsupervised Learning: Introduction

# About the Instructors

Genevera Allen:

- Rice University - Departments of Electrical and Computer Engineering, Stat and CS & Baylor College of Medicine - Neurological Research Institute.
- Founder, Rice D2K Lab
- Research:
  - ▶ Statistical Machine Learning, Data Integration, Graphical Models, Modern Multivariate Analysis, Interpretability & Fairness, Neuroscience, Genomics.

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# About the Instructors

Yufeng Liu:

- University of North Carolina, Chapel Hill - Departments of Statistics and Operations Research, Genetics & Biostatistics.
- Research:
  - ▶ Statistical Machine Learning and Data Mining; High-dimensional Data Analysis; Nonparametric Statistics and Functional Estimation; Personalized Medicine; Bioinformatics.

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# Teaching Assistant

Lili Zheng:

- Postdoctoral Fellow, Rice University.
- PhD Statistics, University of Wisconsin, Madison.
- Research:
  - ▶ Graphical models, missing data, distribution-free inference, tensor data analysis, network Granger causality, dependent data, high-dimensional statistics, stochastic algorithms, and non-convex optimization.

`https://lili-zheng-stat.github.io/`

# Statistical Machine Learning

- “Learn” from current data to make predictions about the future.

Examples?

- Intersection of: Computer Science, Statistics, Applied Math.

# Big Data

Big Data - BIG in Volume, Variety and/or Velocity (or Complexity!).

Common Big Data themes in Statistical Learning:

- Big  $n$ . Large number of observations.
  - ▶ Examples: Internet data, financial transactions, climate data, etc.
- Big  $p$ . Large number of features relative to observations. (High-dimensional data).
  - ▶ Examples: Medical data - genomics, neuroimaging, medical imaging, etc.

# Big Biomedical Data

## Examples:

- High-throughput Genomics (“Omics”).
  - ▶ RNA-sequencing, microarrays, methylation arrays, CGH-arrays, exome sequencing, mass spectrometry, NMR spectroscopy, etc.
- Neuroimaging / neural recordings.
  - ▶ MRI, Functional MRI (fMRI), EEG, MEG, DTI, ECoG, PET, etc.
- Electronic Health Records.
- Medical Imaging.
- Text Data - Pubmed abstracts.

# Data Matrix

Data Matrix:

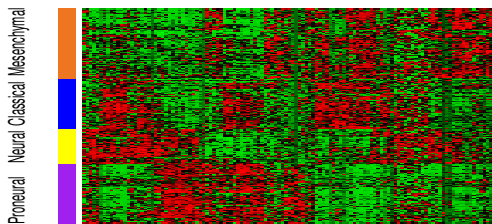
$$\mathbf{X}_{n \times p} = \begin{pmatrix} x_{11} & x_{12} & \dots & x_{1p} \\ \vdots & & \ddots & \\ x_{n1} & x_{n2} & \dots & x_{np} \end{pmatrix}$$

- Rows:  $n$  observations / samples / subjects.
- Columns:  $p$  features / variables.



# Data Matrix

## Example: Omics Data



### Gene Expression Data (Microarray)

- Rows (observations): Subjects ( $n \approx 100 - 500$ ).
- Columns (features): Genes ( $p \approx 500 - 20,000$ ).
- Measurement: Gene expression levels (loosely, how much a gene is turned off or on in a sample).

# Data Matrix

## Example: Text Mining

	data	R	big	cluster	shiny	fast	plot
doc 1	57	1	43	2	0	22	4
doc 2	17	29	2	3	35	6	44
doc 3	47	33	0	0	24	3	19
doc 4	23	0	0	31	0	7	2
doc 5	40	5	28	9	0	21	6
doc 6	8	10	7	46	12	17	9

(Bag-of-Words Format)

- Rows (observations): Documents ( $n \approx 500 - 100,000$ ).
- Columns (features): Words ( $n \approx 100 - 50,000$ ).
- Measurement: Count of how many times words appeared in documents.

# Data Matrix

Example: Image Data



(Handwritten Digits Data)

- Rows (observations): Digits ( $n \approx 10,000$ ).
- Columns (features): Pixels ( $p = 256$ ).
  - ▶ Each digit image is converted to a  $16 \times 16$  grayscale image. The 256 total pixels are vectorized to form the features.
- Measurement: Normalized grayscale intensity of each pixel.

# Unsupervised vs. Supervised Learning

$$\mathbf{X}_{n \times p} = \begin{pmatrix} x_{11} & x_{12} & \dots & x_{1p} \\ \vdots & & \ddots & \\ x_{n1} & x_{n2} & \dots & x_{np} \end{pmatrix}$$

- Rows:  $n$  observations / samples / subjects.
- Columns:  $p$  features / variables.

## Supervised Learning:

$$\mathbf{y} = (y_1, y_2, \dots, y_n)^T$$

- $\mathbf{y}$  -  $n$  labels / outcomes associated with each observation.

## Unsupervised Learning: No outcomes / labels!

# Supervised Learning

## Main Goal

### Prediction!

- Given:  $(Y_n^{train}, \mathbf{X}_{n \times p}^{train})$  (Training Data).
- Training: Use training data to find  $\hat{f}()$  that maps  $\mathbf{X}$  to  $Y$ :  
 $Y = f(\mathbf{X}) + \epsilon$ .
- Prediction: Given new  $\mathbf{X}_{m \times p}^{test}$ , predict  $Y_{m \times 1}^{test}$ :  $\hat{Y}^{test} = \hat{f}(\mathbf{X}^{test})$ .

Examples?

Secondary Goals:

- Feature Selection - What features are associated with the outcome?
- Others?

# Unsupervised Learning

No labels! What is the goal?

## Main Goal

Find some **structure** that characterizes the data.

(Or, find structure in training data that we expect to be present in future data.)

- Find patterns. (PCA, ICA, NMF, MDS)
- Dimension reduction. (PCA)
- Group observations / Group features / Group both. (Clustering)
- Find associations / relationships between features or observations. (Graphical or Network Models)
- Filter features. (Association testing)

# Unsupervised Learning

## Challenges:

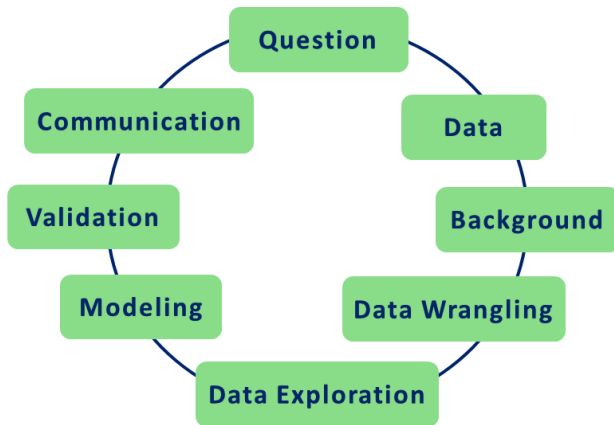
- Difficult to validate unsupervised learning results.
- No validation or test labels to measure prediction accuracy.
- What is meaningful structure in data?

## Uses:

- Data pre-processing / compression / denoising.
- Exploratory data analysis.
  - ▶ Need to use multiple unsupervised learning techniques as each gives slightly different “insights” into data.
- Data visualization.
- Data-Driven Discovery.

# Unsupervised Learning

How does it fit into a data science pipeline?





# Unsupervised Learning

How is it used in Big Biomedical Data?

Case Study: BRCA gene expression data.

- Data Visualization.
  - ▶ Cluster heatmap, graphical models, MDS, PCA.
- Exploratory Analysis.
  - ▶ Clustering / dimension reduction to find cancer subtypes.
- Gene Selection.
  - ▶ Large-scale hypothesis testing to find genes associated with subtypes.
- Gene Interactions.
  - ▶ Graphical models.

# Breakout Discussion

- How will you use Unsupervised Learning?
- What type of big data do you work with?
- What do you hope to learn from this course?

# This Course

## Day 1:

- ① Lecture 1: 11:30-12:20pm - Intro
- ② Lecture 2: 12:30-1:20pm - Dimension Reduction I
- ③ Lecture 3: 1:30-2:20pm - Dimension Reduction II/Lab Intro

*\*All times Pacific.*

# This Course

## Day 2:

- ① Lecture 1: 8-8:50am - Dimension Reduction III / Lab
- ② Lecture 2: 9-9:50am - Dimension Reduction Lab
- ③ Lecture 3: 10-10:50am - Clustering I
- Break
- ④ Lecture 4: 11:30-12:20pm - Clustering II
- ⑤ Lecture 5: 12:30-1:20pm - Clustering III / Lab
- ⑥ Lecture 6: 1:30-2:20pm - Clustering Lab

*\*All times Pacific.*

# This Course

## Day 3:

- ① Lecture 1: 8-8:50am - Testing
- ② Lecture 2: 9-9:50am - Graphical Models I
- ③ Lecture 3: 10-10:50am - Graphical Models II
- Break
- ④ Lecture 4: 11:30-12:20pm - Validation + Final Lab
- ⑤ Lecture 5: 12:30-1:20pm - Final Lab
- ⑥ Lecture 6: 1:30-2:20pm - Final Lab Results + Best Practices

*\*All times Pacific.*