



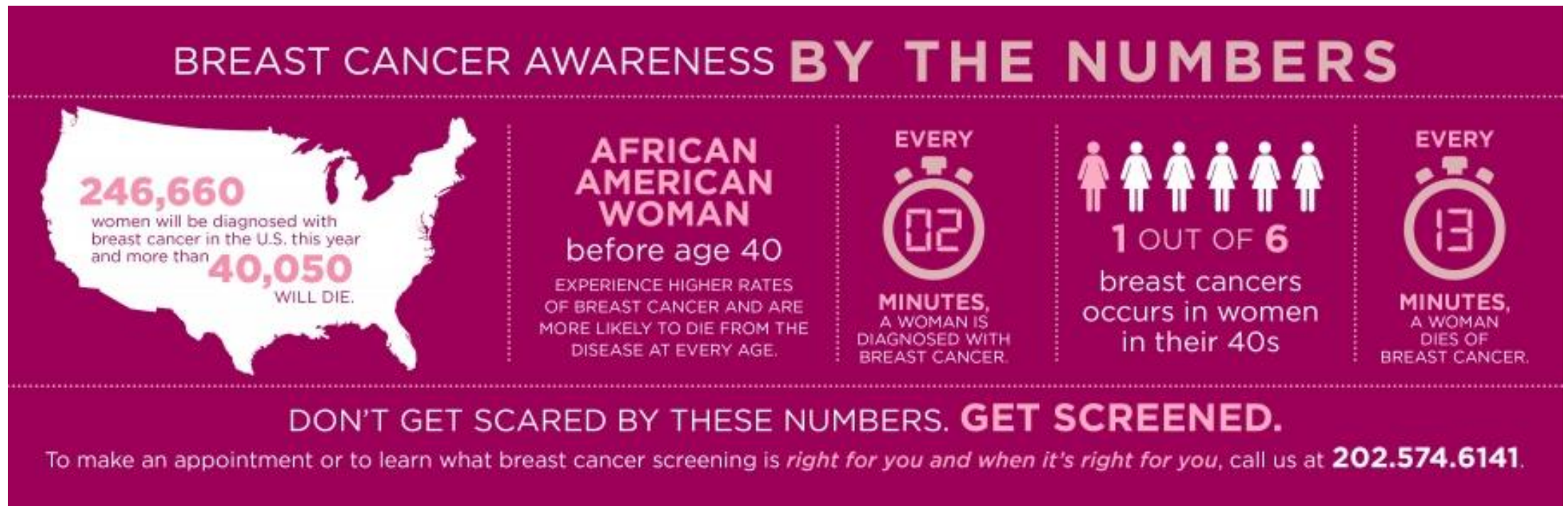
CS+Social Good
CS 106S Winter 2018

today's outline

- 1 breast cancer overview
- 2 intro to machine learning
- 3 cancer diagnosis assignment

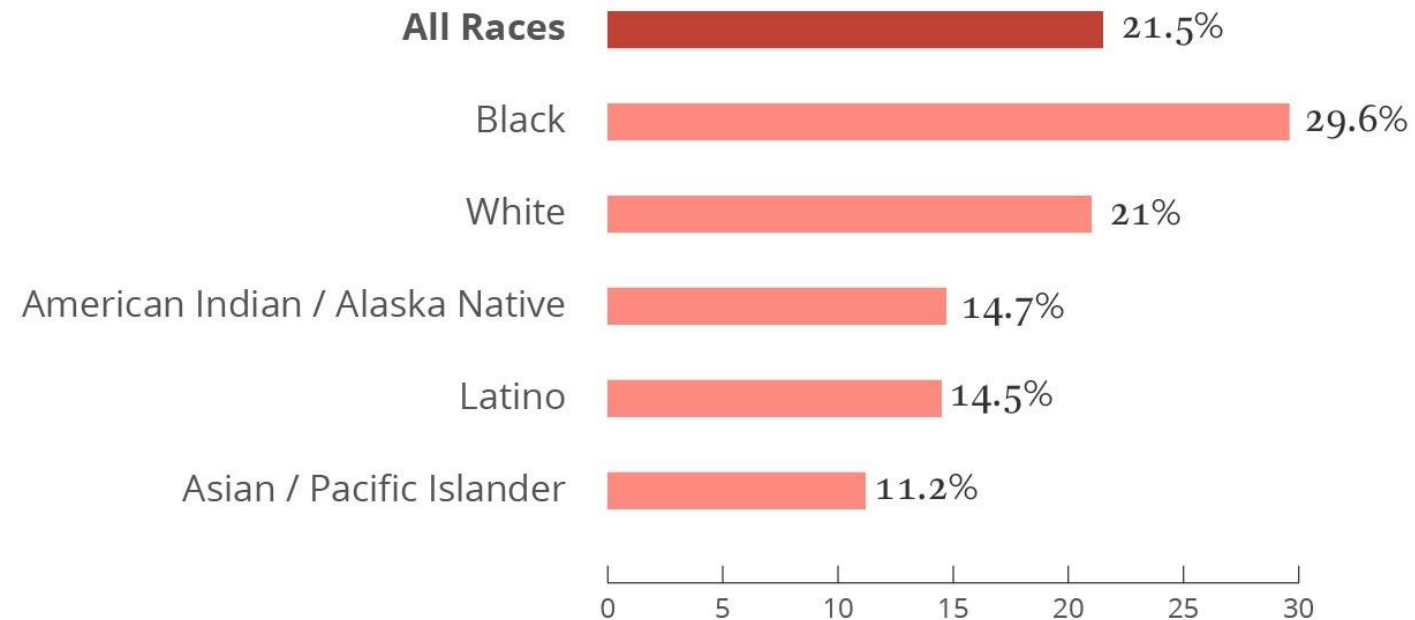


breast cancer



breast cancer

U.S. Breast Cancer Deaths per 100,000 Women by Race/Ethnicity (2009-2013)



breast cancer

- error rates as high as **71%**
- mistakes are common depending on the doctor
- **31%** of all cases misdiagnosed
- **90,000** women misdiagnosed



today's task

Given medical data about cell growths, can we accurately classify tumors as benign or malignant using machine learning?



today's task matters because...

Classifying tumors allows:

- Doctors to diagnose cancer with **greater accuracy**
- Doctors and patients to make **more informed decisions**
- Medical costs to be reduced and diagnosing processes to become **more efficient**



what is machine learning?

Machine learning is a “field of study that gives computers the ability to learn without being explicitly programmed.”

-Arthur Samuel, 1959



types of machine learning

Supervised

Unsupervised



supervised machine learning



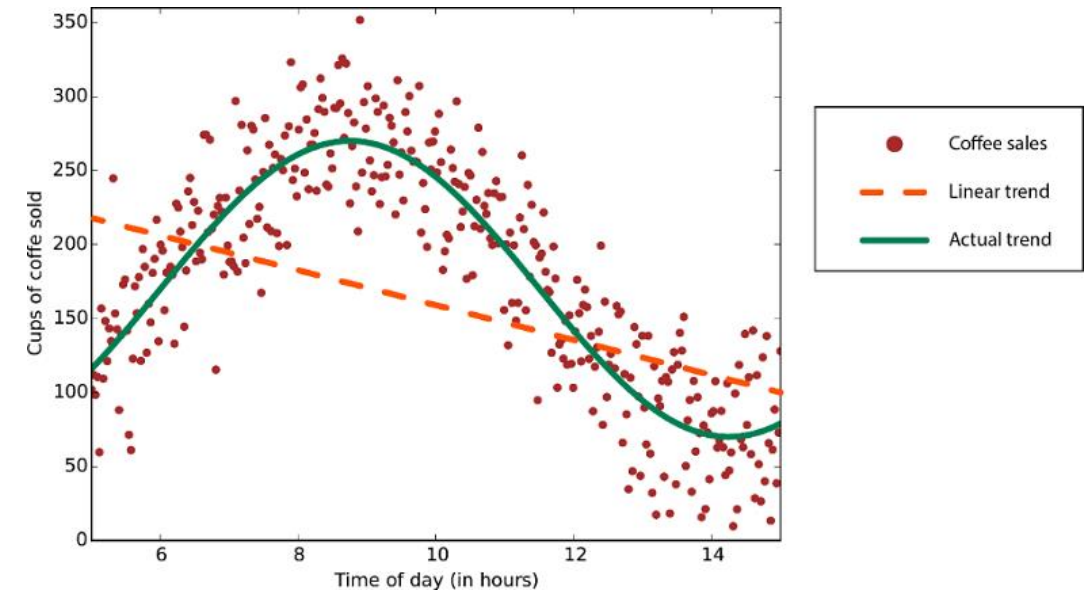
Relationship

“Dog”

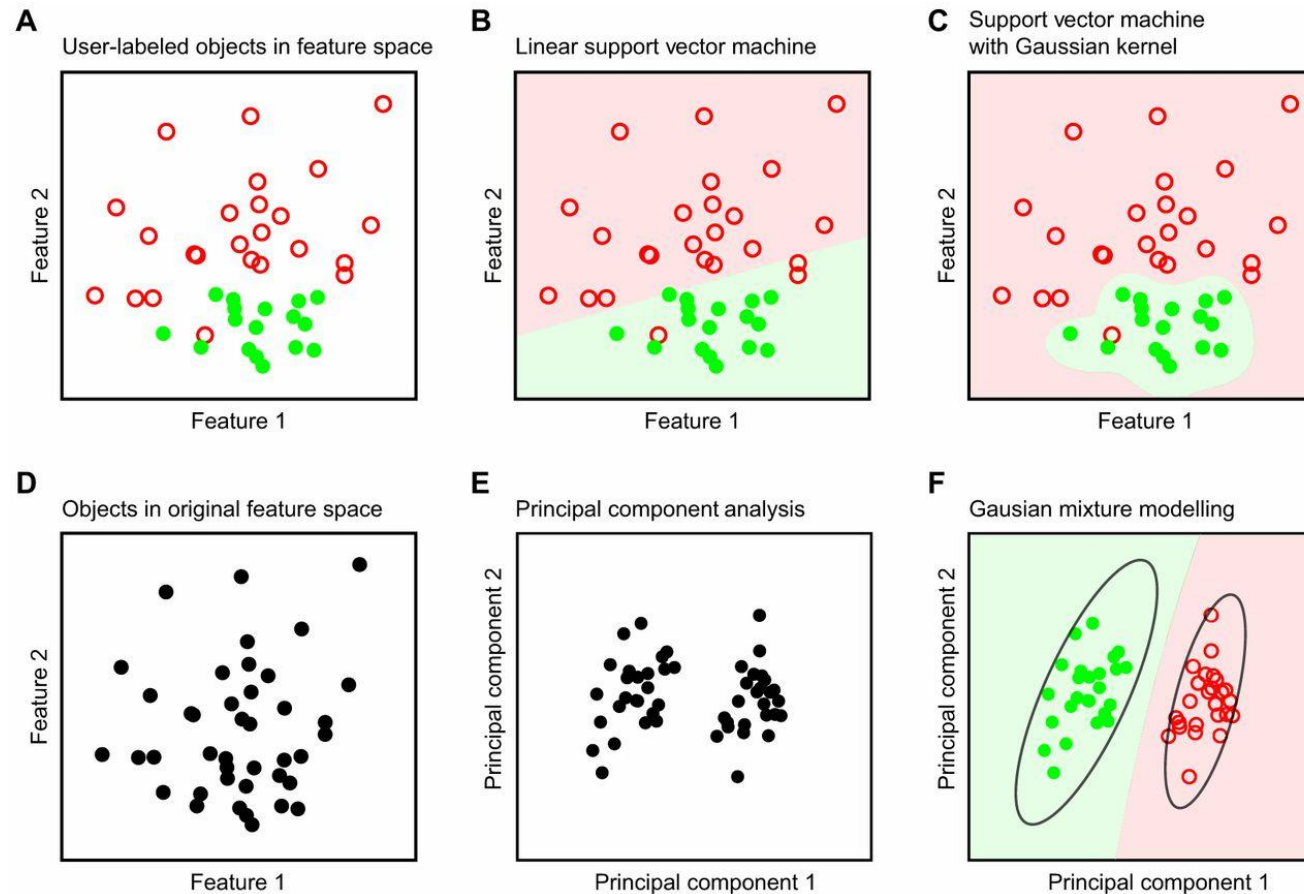


Relationship

“Cat”



unsupervised machine learning



our data

- ~700 total data instances
- We are training on ~630, testing on ~70
- Each data instance has 11 values
 - Sample code number
 - Clump thickness (1-10)
 - Uniformity of Cell Size (1-10)
 - Uniformity of Cell Shape (1-10)
 - Marginal Adhesion (1-10)
 - **Label** (2=benign, 4=malignant)
 - Single Epithelial Cell Size (1-10)
 - Bare Nuclei (1-10)
 - Bland Chromatin (1-10)
 - Normal Nucleoli (1-10)
 - Mitosis (1-10)



implementation steps

- Calculate distance between test instance and every train instance
- Pick the K train instances with the smallest distances
- Of these K train instances, see how many are classified as malignant and how many are classified as benign
- Pick whichever class appears more times as your answer



let's get started!

Build your own breast cancer classifier using JavaScript!

Clone the code at bit.ly/cs106srepo and fill in `cancer-classify.js` located in `cancer-diagnosis/javascript`.

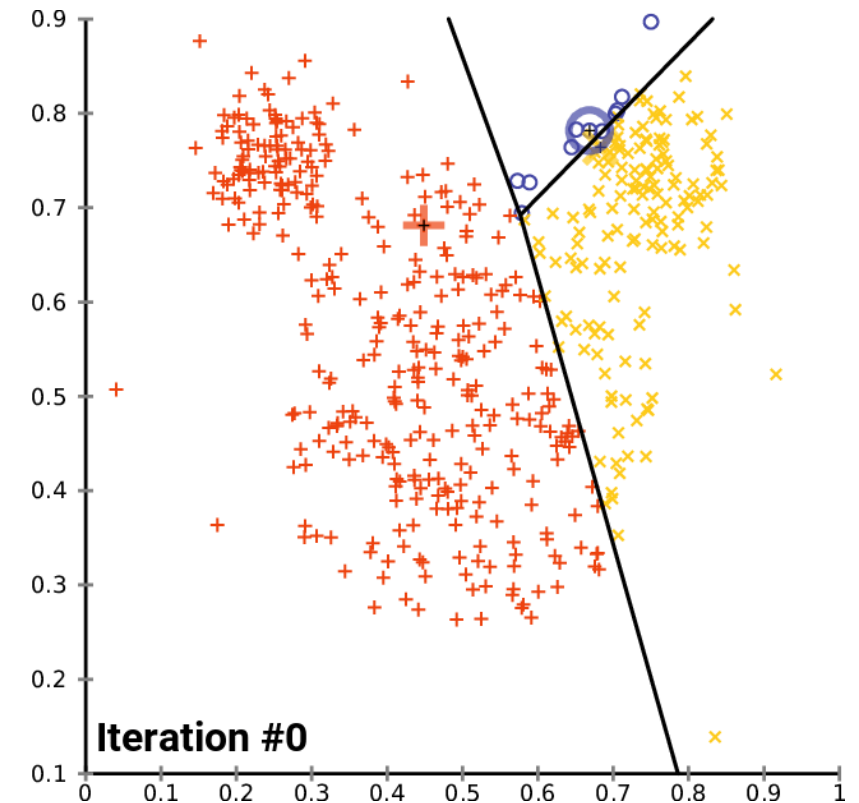
Run `python -m SimpleHTTPServer` in the above folder and examine the console output to test your solution.



extra challenge

Look up k-means and implement it with 2 clusters.

“k-means clustering aims to partition n observations into k clusters in which each observation belongs to the cluster with the nearest mean, serving as a prototype of the cluster.”





See you next time, friends!