

CSC380: Principles of Data Science

Final Project

Kyoungseok Jang

- HW7, final project out
- HW5 solution uploaded

Overview

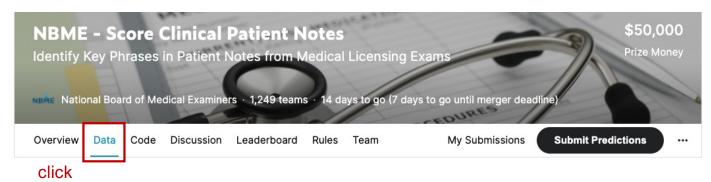
- Participate in NBME task
 - National Board of Medical Examiner
- Mostly guided problem solving, but some open-ended questions
- Extra points for those who achieve high score in the leaderboard.

Problem 1. Familiarize yourself with NBME

- NBME task
 - Clinical skill examination
 - Exam takers interact with patients to write patient note (without missing important information; "features")
 - Want: Automatically grade patient notes! (or, semi-automatically)
- Total 10 types of patients (called "case")

Problem 1. Familiarize yourself with NBME

Understand data



The data:

patient_notes.csv

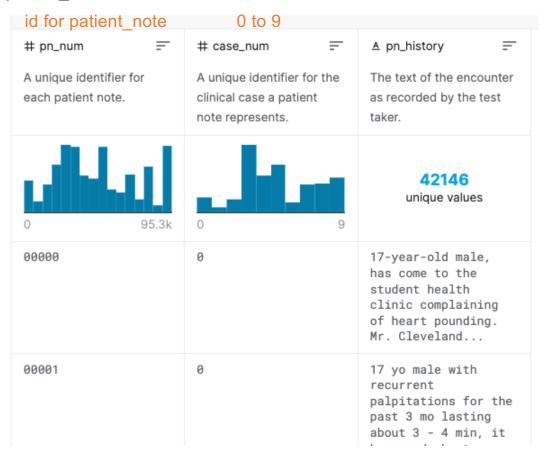
features.csv

For training: train.csv

For submitting your ML code for evaluation: test.csv sample_submission.csv

Problem 1. Familiarize yourself with NBME

patient_notes.csv



Exam taker's answers for each case

patient_notes.csv

Example patient notes

- "17-year-old male, has come to the student health clinic complaining of heart pounding. Mr. Cleveland's mother has given verbal consent for a history, physical examination, and treatment
- -began 2-3 months ago, sudden, intermittent for 2 days (lasting 3-4 min), worsening, non-allev/aggrav
- -associated with dispnea on exersion and rest, stressed out about school
- -reports fe feels like his heart is jumping out of his chest
- -ros:denies chest pain,dyaphoresis,wt loss,chills,fever,nausea,vomiting,pedal edeam
- -pmh:non,meds :aderol (from a friend),nkda
- -fh:father had MI recently,mother has thyroid dz
- -sh:non-smoker,mariguana 5-6 months ago,3 beers on the weekend, basketball at school
- -sh:no std"

17yo male with no pmh here for evaluation of palpitations. States for the last 3-4mo he has felt that his heart with intermittently "beat out of his chest," with some associated difficulty catching his breath. States that the most recent event was 2 days ago, and during activity at a soccer game. He does not seem to note any specific precipitatinig factors at this time. He also states that he feels as if he will faint during these events, but has not lost consciousness at any point. Furthermore, he does endorse theses attacks occuring 1-2 times a month and peak at 4 mins. He denies any stressors at home. ROS: denies weight loss, fevers, recnet illness, change in bowel habits. PMH: negative, PSH negative, FHX mom with thyroid disorder, dad with heart condition and MI at 52yo. SHX no tobacco, ETOH on weekends, Marijuana tried a month ago. Med: is taking some of roommates Adderoll intermittently (last was 2 days ago prior to event). KNDA

patient_notes.csv

"Dillon Cleveland is a 17 year old male with no signflicant past medical history presenting today with ""heart pounding"" for the past 2-3 months. He first noticed an episode when he was sitting down and has had 5-6 over the past 3 months. In the most recent episode has felt light headed and had to sit down while playing basketball. There does not appear to be any precipating factors for there episodes, and has never lost consciousness with them. He has not had any changes in his bowel habits or sleep. No sensitivity to heat or cold. No Weight gain or loss.

Medical history: None Surgical history: None

Medications: Adderal (non-prescription)

Allergies: NKA

Family history: Father had MI 1 year ago. No history of

arrythmias. Mother has a thyroid issue.

SocNo alcohol or tobacco. Tried marijuana once. Drinks one cup of coffee daily. Takes 2x adderal per week for the

previous 8 months."

A variety of writing style!!

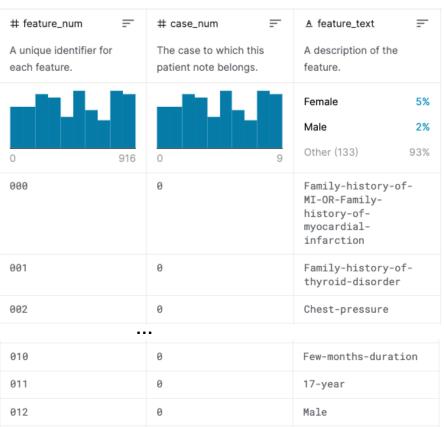
You have a bunch of these for each case_num (=patient).

There are total 10 actual patients.

Note typos

features.csv

Features are not the 'feature vector' we learned! From now on, I will call this 'evaluation' features.



| 101 | 1 | Weight-loss |
|-----|---|---------------------|
| 102 | 1 | Not-sexually-active |

Features: the evaluation target. The first number for the feature_num seems to be case_num

Each case_num has a different number of features.

Visit the Kaggle NBME page to see more on the data

Goal of the competition

The goal of this competition: have a software that, given a patient_note, automatically highlights the relevant features.

Feature 9: Heart pounding

"Dillon Cleveland is a 17 year old male with no signflicant past medical history presenting today with ""heart pounding"" for the past 2-3 months. He first noticed an episode when he was sitting down and has had 5-6 over the past 3 months. In the most recent episode has felt light headed and had to sit down while playing basketball. There does not appear to be any precipating factors for there episodes, and has never lost consciousness with them. He has not had any changes in his bowel habits or sleep. No sensitivity to heat or cold. No Weight gain or loss.

Medical history: None Surgical history: None

Medications: Adderal (non-prescription)

Allergies: NKA

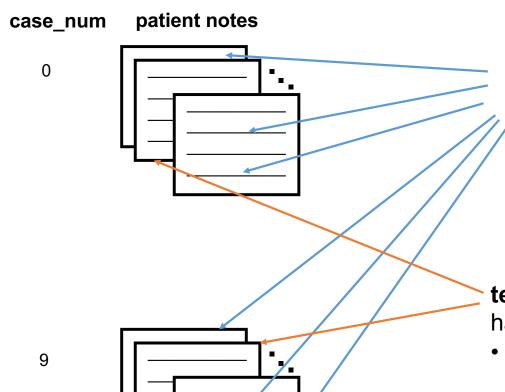
Family history: Father had MI 1 year ago. No history of arrythmias. Mother has a thyroid issue.

SocNo alcohol or tobacco. Tried marijuana once. Drinks one cup of coffee daily. Takes 2x adderal per week for the previous

8 months."

Feature 0: family history of MI

Data Overview



Annotations in **train.csv**: Highlights relavant portion of the note for each evaluation features.

test.csv points out which ones you will have to make prediction

- You will have to highlight portions of texts for each evaluation feature.
- Of course, these never appear in train.csv!

- train.csv contains those annotations for a <u>subset</u> of patient notes.
- Many columns, but the key is

(case_num, patient note number, evaluation feature num, locations)

E.g., (0, 16, 4, ['321 329', '404 413', '652 661'])

Other columns are not essential; just for your convenience.

train.csv

So, train.csv contains those annotations for a <u>subset</u> of patient notes.

| # case_num | = | # pn_num | = | # feature_num | = | ▲ annotation | = | ▲ location | = | |
|---------------------------------------|---|---------------------------------------|-------|--------------------------------|-------|---|--------------------------|--|---|--|
| The case to which patient note belong | | The patient note annotated in this ro | ow. | The feature annotate this row. | ed in | The text(s) within a patient note indicate feature. A feature indicated multiple to within a single note. | ting a nay be imes | Character spar indicating the l each annotation the note. | ocation of | |
| | | | 41 | | | [] | 31% | П | 31% | |
| | | بالمانيال | | | | ['F'] | 2% | ['0 5'] | 1% | |
| 0 | 9 | 16 | 95.3k | 0 | 916 | Other (9597) | 67% | Other (9719) | 68% | |
| 0 | | 00016 | | 000 | | ['dad with recent heart attcak'] ['mom with "thyroid disease'] | | ['696 724'] | python indexing! ('start end' means from start to end-1 | |
| 0 | | 00016 | | 001 | | | | ['668 693'] | | |
| 0 | | 00016 | | 002 | | ['chest pressur | e'] | ['203 217'] | | |
| 0 | | 00016 | | 003 | | <pre>['intermittent episodes', 'episode']</pre> | | ['70 91', '176 183'] | | |

train.csv 14

Be aware:

| # pn_num | = | # feature_num | = | ▲ annotation | = | A location | = |
|---|---|--------------------------------|------|--|------------------------|--|---|
| The patient note annotated in this row. | | The feature annotate this row. | d in | The text(s) within a patient note indicate feature. A feature mindicated multiple ti within a single note. | ing a nay be mes | Character spans indicating the local each annotation withe note. | |
| 00082 | | 009 | | ['heart pounding', 'heart pounding | | ['85 99', '126 '126 131;143 1 | |

some annotations are not contiguous.

E.g.) Heart (.....) pounding

some even overlap with other annotation.

So, you need to use ML to learn from train.csv and be able to mark 'location'.

[Prediction task]

- Given: (case_num, pn_num)
- For each evaluation feature f
 - For each feature you need to perform prediction of those 'locations' for each feature {0,...,12} for case_num=0

[Training]

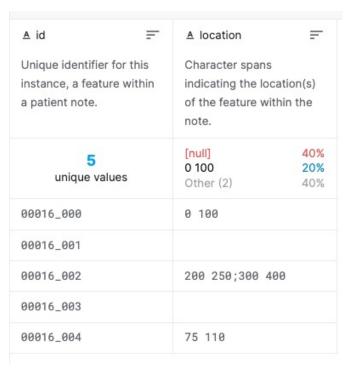
- For each case_num $c \in \{0, ..., 9\}$
 - Use the annotations in train.csv to train a function $g_c(pn_history)$ that returns a list of locations, each corresponding to an evaluation feature.

This will be replaced to a large number of other entries when you submit your answer to Kaggle!

| ∆ id = | # case_num = | # pn_num = | # feature_num = |
|--------------------|-------------------|-------------------|-------------------|
| 5 unique values | 5 total values | 5 total values | 5 total values |
| 00016_000 | 0 | 00016 | 000 |
| 00016_001 | 0 | 00016 | 001 |
| 00016_002 | 0 | 00016 | 002 |
| 00016_003 | 0 | 00016 | 003 |
| 00016_004 | 0 | 00016 | 004 |

sample_submission.csv

Your code will have to write answers like this:



each row corresponds to each row in test.csv

Evaluation criteria

Example

Suppose we have an instance:

These spans give the sets of indices:

We therefore compute:

- TP = size of $\{2, 3, 4\} = 3$
- $FN = size of \{0, 1\} = 2$
- FP = size of $\{7, 8\} = 2$

Repeat for all instances, collect the TPs, FNs, and FPs, and compute the final F1 score.

First, sum up all the TP, FN, and FP computed from every row of submission.csv

Call these summed values as TPs, FNs, and FPs.

```
precision = TPs / (TPs + FPs)
recall = TPs / (TPs + FNs)
```

This is called micro-F1 score!

Example: "One time durign a baskeball game two days ago light headedness, pressure in the chest, but no fainting"

```
Split into words (record their locations as well): ['one', 'time', ..., 'pressure', 'in', 'the, 'chest', 'catching', ...]
```

Idea: what if we build a classifier at the word level?

For each word location [start, end]

- feature vector: [word length, freq('a'), freq('b'), ..., freq('z'), freq('0'), ..., freq('9')]
 e.g., for 'pressure': freq('a') is 0, freq('p') is 1/8, freq('s') is 1/4.
- label: whether the location [start, end] belongs to the annotation.

Note: 'pressure' may appear twice with different label! ('heart' pressure yes, 'air' pressure no)

For making prediction:

- Split by words as before, make prediction for each (+/-)
- Collect the character locations predicted as positive.

Example prediction:

"Most recent episode was accompanied by chest pressure and lasted 10 minutes."

ground truth is "chest pressure" so you get it right except for one space!

Another one:

- "Most recent episode was accompanied by <u>pressure</u> in the <u>chest</u> and lasted 10 minutes."
- Q: 'in' and 'the' received positive label. Why wouldn't it be classified as positive?

How to use ML: Two Words

Observation: Often, words are meaningful 'phrase-wise'.

Idea: Let's consider two words at a time!

Example: "One time durign a baskeball game two days ago light headedness, pressure in the chest, but no fainting"

Extract (and record their locations):

```
['one time', 'time durign', ..., 'headedness, pressure', 'pressure in', 'in the', 'the chest', chest, but', ...]
```

For each two words location [start, end]

- feature vector: [n_characters, freq('a'), freq('b'), ..., freq('z'), freq('0'), ..., freq('9')]
- label: whether the location [start, end] entirely belongs to the annotation.

How to use ML: Two Words

For making prediction:

- Split by two words as before, make classification
- Collect the character locations predicted as positive

// locations may overlap => just take 'logical or'

Example prediction:

"Most recent episode was accompanied by <u>pressure in the chest</u> and lasted 10 minutes."

- 'pressure in' => +
- 'in the' => -
- 'the chest' => +

But some expressions may be meaningful by just one word... what would you do?

How to use ML: Combined

Idea: let's use both 'one word' and 'two words' classifiers!

- Option 1: Build two separate classifiers
- Option 2: Build one classifier How come?

```
Recall that the feature vector is [n_characters, freq('a'), freq('b'), ..., freq('c'), freq('0'), ..., freq('9')]..! 

⇒ Pool all one-word-based data points and two-word-based ones, and train!
```

```
In fact, I would add the number of words as a feature as well:

[n_words, n_characters, freq('a'), freq('b'), ..., freq('z'), freq('0'), ..., freq('9')]
```

I like option 2 better: if n_words mattered, the classifier will pick up that information. Otherwise, it wouldn't.

• E.g., if 'to me' and 'tome' have a different labels, decision tree would use n_words to distinguish it! Otherwise, the decision tree may not even use n_words.

How to use ML: Combined

But how to make predictions?

Given a text

- extract all {(location, word)}, and {(location, two words)}
- make predictions
- collect all positive locations, take 'logical or' (i.e., union of all positive locations)

E.g., if [100,110] and [105,120], then output [100,120]

How to use ML: Combined

But why stop at two words?

In my code provided to you, I use up to 5 words.

- There is a parameter 'W' in the code set to 5.
- Feel free to change it around.

Note that there are many other choices to make!

• E.g., which characters to take into account // rules to separate words (e.g., 'a/b' one word or two words?) // 'stemming' // ...



Useful python functions

eval(expression[, globals[, locals]])

The arguments are a string and optional globals and locals. If provided, *globals* must be a dictionary. If provided, *locals* can be any mapping object.

The *expression* argument is parsed and evaluated as a Python expression (technically speaking, a condition list) using the *globals* and *locals* dictionaries as global and local namespace. If the *globals* dictionary is present and does not contain a value for the key __builtins__, a reference to the dictionary of the built-in module builtins is inserted under that key before *expression* is parsed. That way you can control what builtins are available to the executed code by inserting your own __builtins__ dictionary into *globals* before passing it to eval(). If the *locals* dictionary is omitted it defaults to the *globals* dictionary. If both dictionaries are omitted, the expression is executed with the *globals* and *locals* in the environment where eval() is called. Note, *eval()* does not have access to the nested scopes (non-locals) in the enclosing environment.

The return value is the result of the evaluated expression. Syntax errors are reported as exceptions. Example:

```
>>> x = 1
>>> eval('x+1')
2
```

Note that we use micro F1 score.

In this case, averaging precision/recall may not be a reasonable thing to do. The # of TPs,FPs,FNs could be quite different across the folds.

E.g., precision 1: 1/(1+5) = 1/6, precision 2: 10/(10+2) = 5/6 average: 0.5

micro precision: 11/(11 + 7) = 0.61

In this case, here is a preferred way to compute precision using cross validation.

Gather TP/FP/FN for each fold, using cross validation.

Add up TP/FP/FN and then compute precision

(do the same for recall)

This is what the provided code does.



CSC380: Principles of Data Science

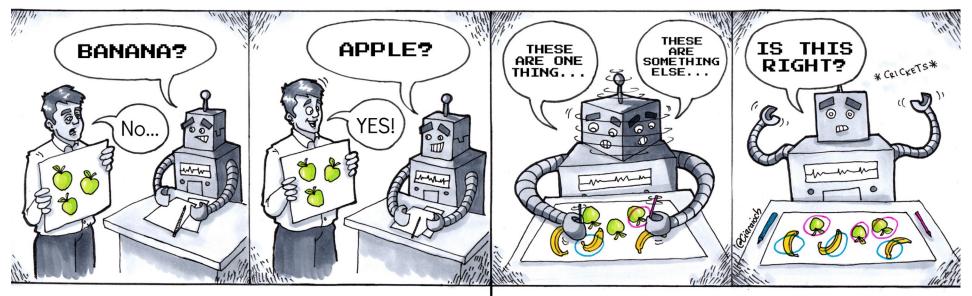
k-means Clustering

Kyoungseok Jang

Slides are in part from Enfa George (TA in 2021)

What is unsupervised learning?

- Learning with unlabeled data
- What can we expect to learn?
 - <u>Clustering</u>: obtain partition of the data that are well-separated.
 - can be viewed as a preliminary classification without predefined class labels.
 - <u>Components</u>: extract common components that compose data points.
 - e.g., topic modeling given a set of articles: each article talks about a few topics => extract the topics that appear frequently.
- Use
 - As a summary of the data
 - Exploratory data analysis: what are the patterns we can get even without labels?
 - Often used as a 'preprocessing techniques'
 - e.g., extract useful **features** using soft clustering assignments (e.g., "gaussian mixture model")



Supervised Learning

Unsupervised Learning



Task 1: Group These Set of Document into 3 Groups based on meaning

Doc1: Health, Medicine, Doctor

Doc 2 : Machine Learning, Computer

Doc 3 : Environment, Planet

Doc 4: Pollution, Climate Crisis

Doc 5: Covid, Health, Doctor



Task 1: Group These Set of Document into 3 Groups based on meaning

Doc1: Health, Medicine, Doctor

Doc 2: Machine Learning, Computer

Doc 3: Environment, Planet

Doc 4: Pollution, Climate Crisis

Doc 5: Covid, Health, Doctor



Task 1: Group These Set of Document into 3 Groups based on meaning

Doc1: Health, Medicine, Doctor

Doc 5: Covid, Health, Doctor

Doc 3 : Environment,

Planet

Doc 4 : Pollution, Climate

Crisis

Doc 2 : Machine Learning, Computer



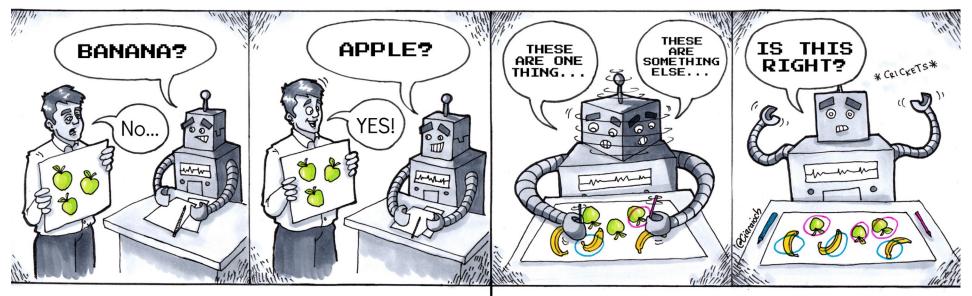
Task 2: Recommendation

- Discover the probability of the co-occurrence of items in a collection
 - Market basket analysis
 - Semantic clustering (Topic modeling)
 - Movie recommendation





From: https://www.simplilearn.com/tutorials/machine-learning-tutorial/supervised-and-unsupervised-learning
And https://developers.google.com/machine-learning/recommendation/collaborative/basics



Supervised Learning

Unsupervised Learning

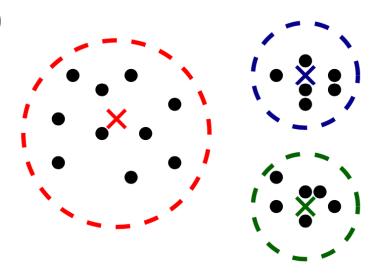


Clustering

• Input: k: the number of clusters (hyperparameter)

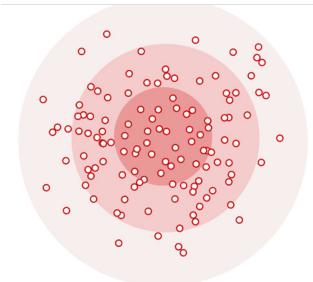
$$S = \{x_1, \dots, x_n\}$$

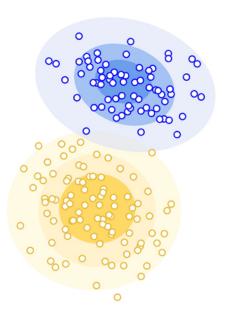
- Output
 - partition $\{G_i\}_{i=1}^k$ s.t. $S = \bigcup_i G_i$ (disjoint union).
 - often, we also obtain 'centroids'



Distribution-based Clustering

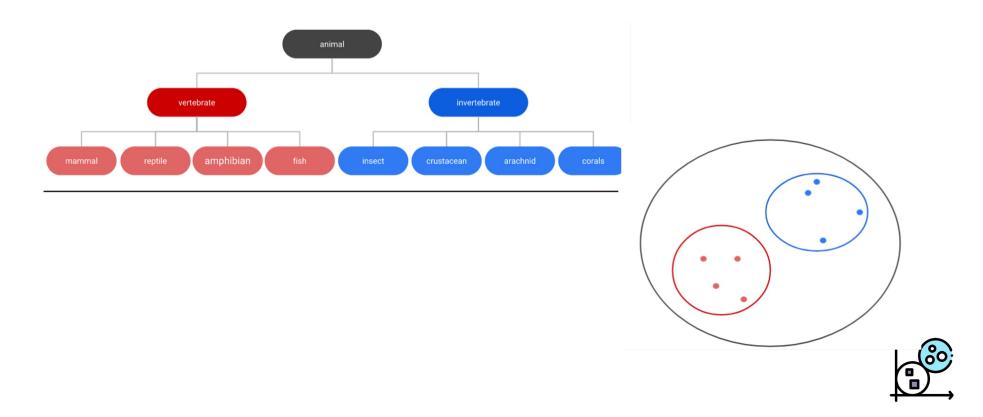
(probabilistic treatment)



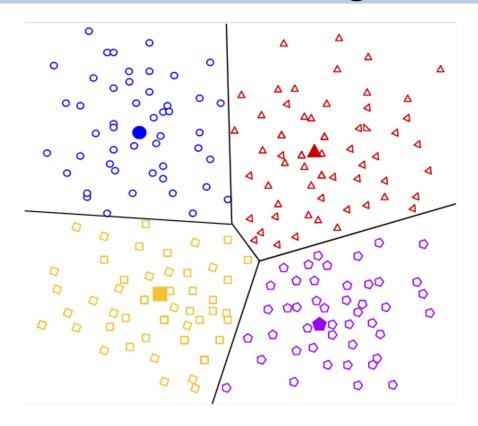




Hierarchical Clustering



Centroid-based Clustering

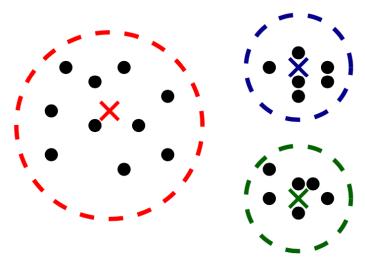




• Input: *k*: the number of clusters (hyperparameter)

$$S = \{x_1, \dots, x_n\}$$

- Output
 - partition $\{G_i\}_{i=1}^k$ s.t. $S = \bigcup_i G_i$ (disjoint union).
 - · often, we also obtain 'centroids'



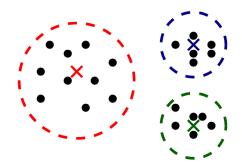
- Q: if we are given the groups, what would be a reasonable definition of centroids?
 - The **point** that has the minimum average **distance** to the datapoints?
 - The **datapoint** that has the minimum average **distance** to the datapoints?
 - The **point** that has the minimum average **squared distance** to the datapoints?

=> Turns out, the last one corresponds to the average point!

k-means clustering

• Idea: if someone gave us k reasonable centroids c_1, \dots, c_k , we can partition the data with them.

$$A(x) = \arg\min_{j \in [k]} ||x - c_j||_2^2$$



 But we don't have those centroids ⇒ Let's find them with an optimization formulation.

$$\arg \min_{c_1, \dots, c_k} \sum_{i=1}^n ||x_i - A(x_i)||_2^2 = \arg \min_{c_1, \dots, c_k} \sum_{i=1}^n \min_{j \in [k]} ||x_i - c_j||_2^2$$

=> NP-hard

k-means Clustering

Lloyd's algorithm: solve it approximately (heuristic)

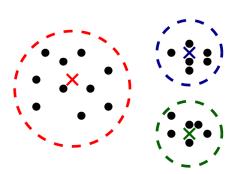
(but people just say it is kmeans clustering algorithm)

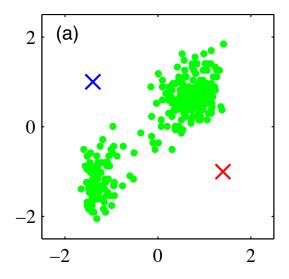
(1957 by Stuart P. Lloyd but independently developed by Joel Max in 1960)



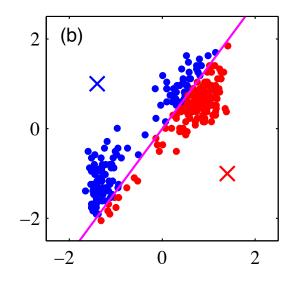
- If you knew the cluster assignments... just find the centroids as the average
- If you knew the centroids... make cluster assignments by the closest centroid.

Why not: start from some centroids and then alternate between the two?

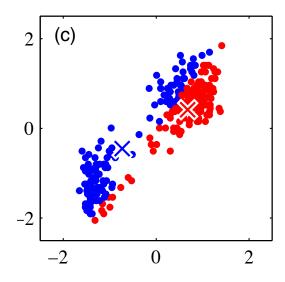




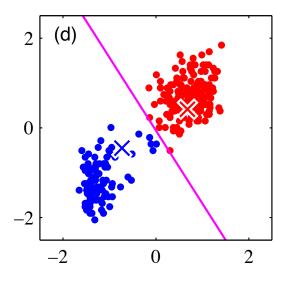
Arbitrary/random initialization of c_1 and c_2



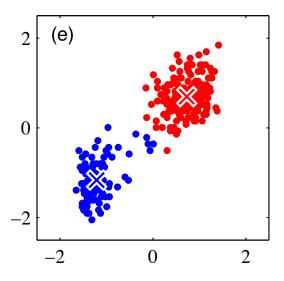
(A) update the cluster assignments.



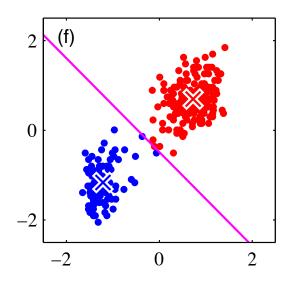
(B) Update the centroids $\{c_j\}$



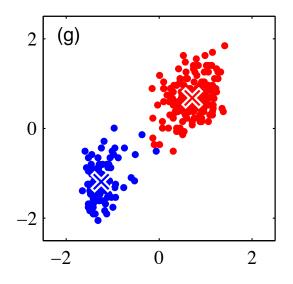
(A) update the cluster assignments.



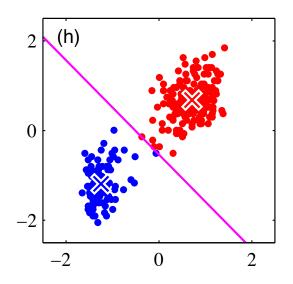
(B) Update the centroids $\{c_j\}$



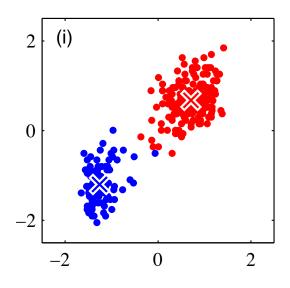
(A) update the cluster assignments.



(B) Update the centroids $\{c_j\}$

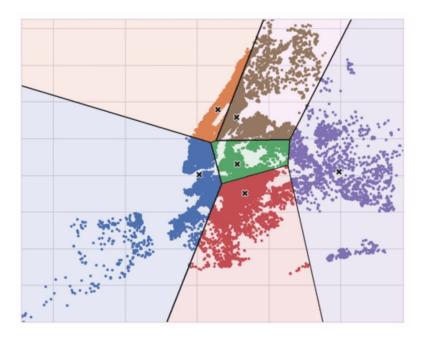


(A) update the cluster assignments.



(B) Update the centroids $\{c_j\}$

Iterating until Convergence





Input: k: num. of clusters, $S = \{x_1, ..., x_n\}$

[Initialize] Pick $c_1, ..., c_k$ as randomly selected points from S (see next slides for alternatives)

For t=1,2,...,max_iter

- [Assignments] $\forall x \in S$, $a_t(x) = \arg\min_{j \in [k]} ||x c_j||_2^2$
- If $t \neq 1$ AND $a_t(x) = a_{t-1}(x), \forall x \in S$
 - break
- [Centroids] $\forall j \in [k], c_j \leftarrow \text{average}(\{x \in S: a_t(x) = j\})$

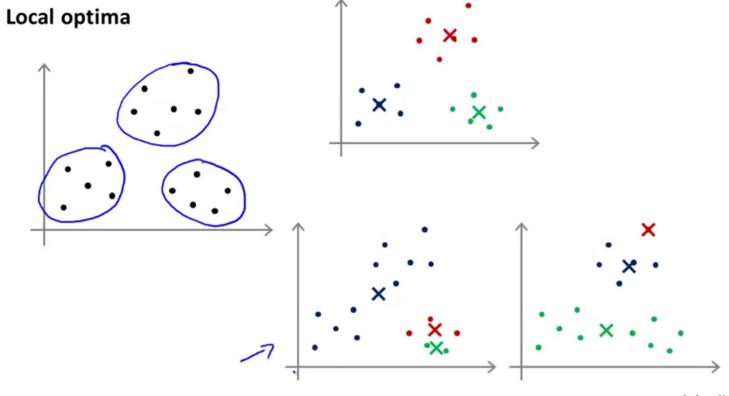
Output: $c_1, ..., c_k$ and $\{a_t(x_i)\}_{i \in [n]}$

But,

It may converge to a local rather than global minimum.

$$\arg \min_{c_1, \dots, c_k} \sum_{i=1}^n \min_{j \in [k]} ||x_i - c_j||_2^2$$









Issue 1: Unreliable solution

- You usually get suboptimal solutions
- You usually get different solutions every time you run.
- <u>Standard practice</u>: Run it 50 times and take the one that achieves the smallest objective function
 - Recall: $\min_{c_1,...,c_k} \sum_{i=1}^n \min_{j \in [k]} ||x_i c_j||_2^2$

Each run of algorithm outputs $c_1, ..., c_k$. Compute this to evaluate the quality!

- And/or, change the initialization (next slide)
 - Idea: ensure that we pick a widespread c_1, \dots, c_k

Two alternative initializations.

- Farthest-first traversal \Rightarrow Sequentially choose c_j that are the farthest from the previously-chosen.
 - Pick $c_1 \in \{x_1, ..., x_n\}$ arbitrarily (or randomly)
 - For j = 2, ..., k
 - Pick $c_j \in \mathbb{R}^d$ as a point in $\{x_1, \dots, x_n\}$ that maximizes the squared distances to c_1, \dots, c_{j-1} .

$$c_j = \arg\max_{i \in [n]} \min_{j'=1,\dots,j-1} ||x_i - c_{j'}||_2^2$$

• k-means++

- Pick $c_1 \in \{x_1, ..., x_n\}$ uniformly at random
- For j = 2, ..., k
 - Define a distribution $\forall i \in [n]$, $\mathbb{P}(c_j = x_i) \propto \min_{j' = 1, \dots, j 1} \|x_i c_{j'}\|_2^2$ More likely to choose x_i

• Draw c_j from the distribution above.

More likely to choose x_i that is farthest from already-chosen centroids.

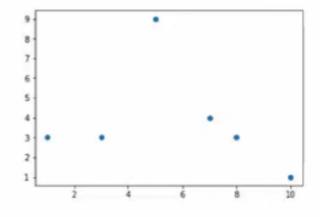
=> has a mathematical guarantee that it will be better than an arbitrary starting point!

Suppose we have the small dataset

(7,4),(8,3),(5,9),(3,3),(1,3),(10,1) to which we wish to assign 3 clusters.

We begin by randomly selecting (7,4) to be a cluster center.

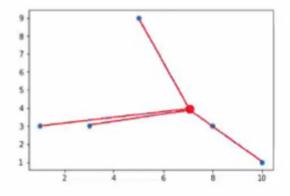
| X | $\min(d(x,z_i)^2)$ |
|--------|--------------------|
| (7,4) | |
| (8,3) | |
| (5,9) | |
| (3,3) | |
| (1,3) | |
| (10,1) | |





We begin by randomly selecting (7,4) to be a cluster center.

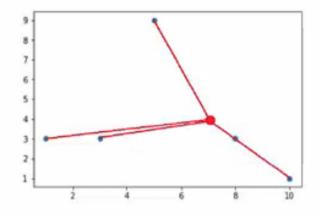
| X | $\min(d(x,z_i)^2)$ |
|--------|--------------------|
| (7,4) | -, |
| (8,3) | 2 |
| (5,9) | 29 |
| (3,3) | 17 |
| (1,3) | 37 |
| (10,1) | 18 |





We begin by randomly selecting (7,4) to be a cluster center.

| X | prob |
|--------|--------|
| (7,4) | - |
| (8,3) | 2/103 |
| (5,9) | 29 103 |
| (3,3) | 17/103 |
| (1,3) | 37/103 |
| (10,1) | 18/103 |

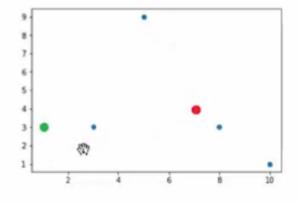




From Sara Jensen's Youtube Channel

We add (1,3) to the list of cluster centers.

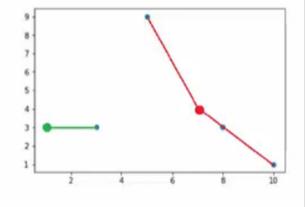
| X | $\min(d(x,z_i)^2)$ |
|--------|--------------------|
| (7,4) | -, |
| (8,3) | |
| (5,9) | |
| (3,3) | 4 100 |
| (1,3) | - |
| (10,1) | * |





We add (1,3) to the list of cluster centers.

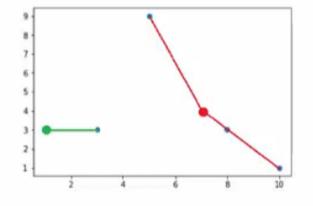
| X | $\min(d(x,z_i)^2)$ |
|--------|--------------------|
| (7,4) | - |
| (8,3) | 2 |
| (5,9) | 29 |
| (3,3) | 4 |
| (1,3) | - |
| (10,1) | 18 |





We add (1,3) to the list of cluster centers.

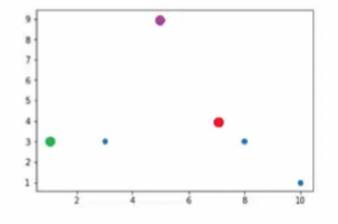
| X | prob |
|--------|-------|
| (7,4) | - |
| (8,3) | 2/55 |
| (5,9) | 29/55 |
| (3,3) | 4/55 |
| (1,3) | - |
| (10,1) | 18/55 |





We add (5,9) to the list of cluster centers.

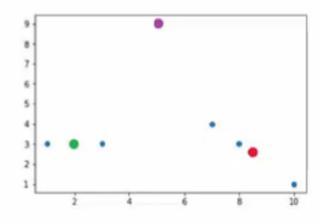
| X | prob |
|--------|------|
| (7,4) | - |
| (8,3) | |
| (5,9) | - |
| (3,3) | |
| (1,3) | - |
| (10,1) | |



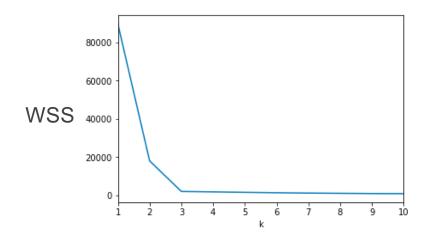


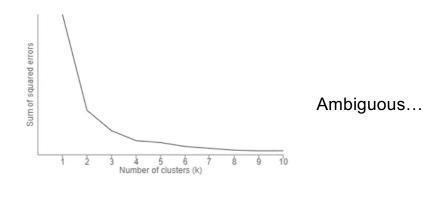
We now run k-means with initialized centers (7,4),(1,3), and (5,9).

| X | prob |
|--------|------|
| (7,4) | - |
| (8,3) | |
| (5,9) | - |
| (3,3) | |
| (1,3) | - |
| (10,1) | |



- No principled way no test set!
- Elbow method: see where you get saturation.
 - When WSS (Within-sum of squares) starts diminishing?





https://medium.com/analytics-vidhya/how-to-determine-the-optimal-k-for-k-means-708505d204eb