

College of Media, Communication and Information



Clustering

INFO-2301: Quantitative Reasoning 2 Michael Paul and Jordan Boyd-Graber SLIDES ADAPTED FROM DAVE BLEI AND LAUREN HANNAH

- What is clustering?
- K-Means
- Gaussian Mixture Models

Clustering



Uses:

- genomics
- medical imaging
- social network analysis
- recommender systems
- market segmentation
- voter analysis

Microarray Gene Expression Data



From: "Skin layer-specific transcriptional profiles in normal and recessive yellow (Mc1re/Mc1re) mice" by April and Barsh in Pigment Cell Research (2006)

Medical Imaging (MRIs and PET scans)



From: "Fluorodeoxyglucose positron emission tomography of mild cognitive impairment with clinical follow-up at 3 years" by Pardo et al. in Alzheimer's and Dementia (2010)

Social Networks



Recommender Systems



From: tech.hulu.com/blog/



Market Segmentation



From: mappinganalytics.com/map/segmentation-maps/segmentation-map.html

Voter Analysis













- soccer moms (female, middle aged, married, middle income, white, kids, suburban)
- Nascar dads (male, middle aged, married, middle income, white, kids, Southern, suburban or rural)
- security moms (...)
- low information voters
- Ivy League Elites





Clustering

Introduction to Data Science University of Colorado Boulder SLIDES ADAPTED FROM LAUREN HANNAH Questions:

- how do we fit clusters?
- how many clusters should we use?
- how should we evaluate model fit?

How do we fit the clusters?

- simplest method: K-means
- requires: real-valued data
- idea:
 - pick K initial cluster means
 - associate all points closest to mean k with cluster k
 - use points in cluster k to update mean for that cluster
 - re-associate points closest to new mean for k with cluster k
 - use new points in cluster k to update mean for that cluster
 - o ...
 - stop when no change between updates

Animation at: http://shabal.in/visuals/kmeans/1.html

Data:

<i>x</i> ₁	<i>x</i> ₂	<i>σ</i> -	•
0.4	-1.0		
-1.0	-2.2	N -	• • • •
-2.4	-2.2	÷ -	-
-1.0	-1.9	ž	°
-0.5	0.6	0 -	-
-0.1	1.7	_	
1.2	3.3	ī -	
3.1	1.6	9 -	•
1.3	1.6		
2.0	0.8		-2 -1 U 1 2 3 X1

K-Means: Example

Pick K centers (randomly):



(-1, -1) and (0, 0)

Calculate distance between points and those centers:

<i>x</i> ₁	<i>x</i> ₂	(-1,-1)	(0,0)
0.4	-1.0	1.4	1.1
-1.0	-2.2	1.2	2.4
-2.4	-2.2	1.9	3.3
-1.0	-1.9	0.9	2.2
-0.5	0.6	1.6	0.8
-0.1	1.7	2.9	1.7
1.2	3.3	4.8	3.5
3.1	1.6	4.8	3.4
1.3	1.6	3.5	2.1
2.0	0.8	3.5	2.2

Choose mean with smaller distance:

<i>x</i> ₁	<i>x</i> ₂	(-1,-1)	(0,0)
0.4	-1.0	1.4	1.1
-1.0	-2.2	1.2	2.4
-2.4	-2.2	1.9	3.3
-1.0	-1.9	0.9	2.2
-0.5	0.6	1.6	0.8
-0.1	1.7	2.9	1.7
1.2	3.3	4.8	3.5
3.1	1.6	4.8	3.4
1.3	1.6	3.5	2.1
2.0	0.8	3.5	2.2

New clusters:



Refit means for each cluster:

- cluster 1: (-1.0, -2.2), (-2.4, -2.2), (-1.0, -1.9)
- new mean: (-1.5, -2.1)
- cluster 2: (0.4, -1.0), (-0.5, 0.6), (-0.1, 1.7), (1.2, 3.3), (3.1, 1.6), (1.3, 1.6), (2.0, 0.8)
- new mean: (1.0, 1.2)



Recalculate distances for each cluster:

<i>x</i> ₁	<i>x</i> ₂	(-1.5, -2.1)	(1.0, 1.2)
0.4	-1.0	2.2	2.3
-1.0	-2.2	0.5	4.0
-2.4	-2.2	1.0	4.9
-1.0	-1.9	0.5	3.8
-0.5	0.6	2.8	1.7
-0.1	1.7	4.1	1.2
1.2	3.3	6.0	2.1
3.1	1.6	5.8	2.0
1.3	1.6	4.6	0.5
2.0	0.8	4.6	1.1

Choose mean with smaller distance:

<i>x</i> ₁	<i>x</i> ₂	(-1.5, -2.1)	(1.0, 1.2)
0.4	-1.0	2.2	2.3
-1.0	-2.2	0.5	4.0
-2.4	-2.2	1.0	4.9
-1.0	-1.9	0.5	3.8
-0.5	0.6	2.8	1.7
-0.1	1.7	4.1	1.2
1.2	3.3	6.0	2.1
3.1	1.6	5.8	2.0
1.3	1.6	4.6	0.5
2.0	0.8	4.6	1.1

New clusters:



Refit means for each cluster:

- cluster 1: (0.4, -1.0), (-1.0, -2.2), (-2.4, -2.2), (-1.0, -1.9)
- new mean: (-1.0, -1.8)
- cluster 2: (-0.5, 0.6), (-0.1, 1.7), (1.2, 3.3), (3.1, 1.6), (1.3, 1.6), (2.0, 0.8)

new mean: (1.2, 1.6)



Recalculate distances for each cluster:

<i>x</i> ₁	<i>x</i> ₂	(-1.0, -1.8)	(1.2, 1.6)
0.4	-1.0	1.6	2.7
-1.0	-2.2	0.4	4.4
-2.4	-2.2	1.5	5.2
-1.0	-1.9	0.1	4.1
-0.5	0.6	2.4	2.0
-0.1	1.7	3.6	1.2
1.2	3.3	5.6	1.7
3.1	1.6	5.3	1.9
1.3	1.6	4.1	0.1
2.0	0.8	4.0	1.2

Select smallest distance and compare these clusters with previous:

<i>x</i> ₁	<i>x</i> ₂	(-1.0, -1.8)	(1.2, 1.6)
0.4	-1.0	1.6	2.7
-1.0	-2.2	0.4	4.4
-2.4	-2.2	1.5	5.2
-1.0	-1.9	0.1	4.1
-0.5	0.6	2.4	2.0
-0.1	1.7	3.6	1.2
1.2	3.3	5.6	1.7
3.1	1.6	5.3	1.9
1.3	1.6	4.1	0.1
2.0	0.8	4.0	1.2

Table: New Clusters

Table: Old Clusters

(-1.5, -2.1)	(1.0, 1.2)
2.2	2.3
0.5	4.0
1.0	4.9
0.5	3.8
2.8	1.7
4.1	1.2
6.0	2.1
5.8	2.0
4.6	0.5
4.6	1.1

K-means can be used for *image* segmentation

- partition image into multiple segments
- find boundaries of objects
- make art





K-Means Clustering

What if our data look like this?



K-Means Clustering

True clustering:



K-Means Clustering

K-means clustering (K = 2):







Clustering

Introduction to Data Science University of Colorado Boulder SLIDES ADAPTED FROM LAUREN HANNAH K-means associates data with cluster centers.

What if we actually modeled the data?

- real-valued data
- observation \mathbf{x}_i in cluster c_i
- have K clusters
- model each cluster with a Gaussian distribution

$$\mathbf{x}_i | c_i = k \sim N(\mu_k, \Sigma_k)$$

• μ_k is mean vector, Σ_k is covariance matrix

Mixture Models

Gaussian mixture model (K = 2):



Why mixture models?

- more flexible: can account for clusters with different shapes
- have data model (will be useful for choosing *K*)
- less sensitive to data scaling

Multivariate Gaussian

Multivariate Gaussian distribution for $\mathbf{x} \in \mathbf{R}^d$:

$$p(\mathbf{x} | \boldsymbol{\mu}, \boldsymbol{\Sigma}) = (2\pi)^{-\frac{d}{2}} |\boldsymbol{\Sigma}|^{-\frac{1}{2}} e^{-\frac{1}{2}(\mathbf{x}-\boldsymbol{\mu})^{\mathsf{T}}\boldsymbol{\Sigma}^{-1}(\mathbf{x}-\boldsymbol{\mu})}$$

- µ is vector of means
- Σ is covariance matrix



Multivariate Gaussian

pdf when
$$\mu = [0, 0]$$
 and $\Sigma = \begin{bmatrix} 0.9 & 0.4 \\ 0.4 & 0.3 \end{bmatrix}$:



Multivariate Gaussian



Mixture model:

- observation x_i in cluster c_i with K clusters
- model each cluster with a Gaussian distribution

$$\mathbf{x}_i | c_i = k \sim N(\mu_k, \Sigma_k)$$

How do we find c_1, \ldots, c_n (clusters) and $(\mu_1, \Sigma_1), \ldots, (\mu_K, \Sigma_K)$ (cluster centers)?

First, let's simplify the model:

· covariance matrices have only diagonal elements,

$$\Sigma = \begin{bmatrix} \sigma_1^2 & 0 & \dots & 0 \\ 0 & \sigma_2^2 & \dots & 0 \\ \dots & \dots & \dots & 0 \\ 0 & 0 & 0 & \sigma_K^2 \end{bmatrix}$$

• set
$$\sigma_1^2 = \cdots = \sigma_K^2$$
, suppose known

Next, use a method similar to K-means:

- start with random cluster centers
- associate observations to clusters by (log-)likelihood,

$$\ell(\mathbf{x}_{i} | c_{i} = k) = -\frac{d}{2} \log(2\pi) - \frac{1}{2} \log\left(\prod_{j=1}^{d} \sigma_{k,j}^{2}\right) - \frac{1}{2} \sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2} / \sigma_{k,j}^{2}$$
$$\propto -d \log(\sigma_{k}) - \frac{1}{2\sigma_{k}^{2}} \sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2}$$
$$\propto -\sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2}$$

• refit centers μ_1, \ldots, μ_K given clusters by

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_i=k} x_{i,j}$$

recluster observations...

clustering with K-means

minimize distance

$$d(\mathbf{x}_{i}, \mu_{k}) = \sqrt{\sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2}}$$

clustering with GMM maximize likelihood

$$\ell(\mathbf{x}_i | c_i = k) \propto -\sum_{j=1}^{a} (x_{i,j} - \mu_{k,j})^2$$

update means with K-means

use average

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_i=k} x_{i,j}$$

update means with GMM use average

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_j = k} x_{i,j}$$

OK, now what if

$$\Sigma = \left[egin{array}{ccccc} \sigma_1^2 & 0 & \dots & 0 \ 0 & \sigma_2^2 & \dots & 0 \ \dots & \dots & \dots & 0 \ 0 & 0 & 0 & \sigma_K^2 \end{array}
ight]$$

and $\sigma_1^2, \ldots, \sigma_K^2$ can take different values?

- use same algorithm
- update μ_k and σ_k^2 with maximum likelihood estimator,

$$\mu_{k,j} = \frac{1}{n_k} \sum_{c_i = k} x_{i,j}$$

$$\sigma_{k,j}^2 = \frac{1}{n_k} \sum_{c_i = k} (x_{i,j} - \mu_{k,j})^2$$

Data:

<i>x</i> ₁	<i>x</i> ₂	o
-3.7	-0.4	ιο –
0.4	0.1	
0.4	-1.7	м –
-0.4	-1.0	°
-1.3	-1.7	- ~ Z
1.0	3.3	
1.2	5.2	° - °
1.3	0.3	- 0 0
1.1	-0.8	
0.5	2.8	∼
		x1

- pick centers and variances, $\mu_1 = [-1, -1]$, $\sigma_1^2 = [1, 1]$, $\mu_1 = [1, 1]$, $\sigma_1^2 = [1, 1]$
- compute (proportional) log likelihoods,

$$\ell(\mathbf{x}_{i} | c_{i} = k) = -\sum_{j=1}^{d} \log(\sigma_{j}) - \frac{1}{2} \sum_{j=1}^{d} (x_{i,j} - \mu_{k,j})^{2} / \sigma_{k,j}^{2}$$

<i>x</i> ₁	<i>x</i> ₂	<i>k</i> = 1	k = 2
-3.7	-0.4	-3.8	-12.1
0.4	0.1	-1.6	-0.6
0.4	-1.7	-1.2	-3.8
-0.4	-1.0	-0.2	-3.0
-1.3	-1.7	-0.3	-6.3
1.0	3.3	-11.2	-2.6
1.2	5.2	-22.0	-9.0
1.3	0.3	-3.6	-0.3
1.1	-0.8	-2.2	-1.6
0.5	2.8	- 8.2	-1.7

fit new means and variances:

$$\mu_1 = [-1.3, -1.2]$$

$$\sigma_1^2 = [3.1, 0.4]$$

$$\mu_2 = [0.9, 1.8]$$

$$\sigma_2^2 = [0.2, 5.4]$$

compute new distances...

<i>x</i> ₁	<i>x</i> ₂	k = 1	k = 2
-3.7	-0.4	-1.8	-70.8
0.4	0.1	-2.7	-1.0
0.4	-1.7	-0.8	-2.0
-0.4	-1.0	-0.3	-6.8
-1.3	-1.7	-0.5	-16.6
1.0	3.3	-27.4	-0.1
1.2	5.2	-55.9	-1.3
1.3	0.3	-4.3	-0.7
1.1	-0.8	-1.2	-0.6
0.5	2.8	-21.3	-0.7

No change, so clusters are final



k-means is fast and simple, but ...

- What if your data are discrete?
- What if each data point has more than one cluster? (digits vs. documents)
- What if you don't know the number of clusters?

- Clustering helps discover patterns
- *k*-means is a simple approach
- Gaussian mixture models more probabilistic foundation