



Clustering

Introduction to Data Science University of Colorado Boulder

SLIDES ADAPTED FROM LAUREN HANNAH

Clustering

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
 - 0
 - stop when no change between updates

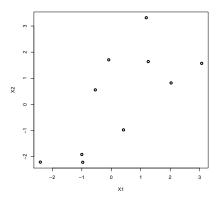
K-Means

Animation at:

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

Data:

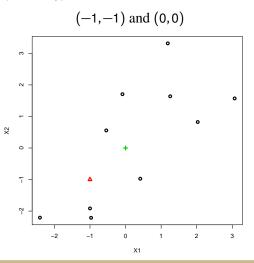
<i>X</i> ₁	<i>X</i> ₂
0.4	-1.0
-1.0	-2.2
-2.4	-2.2
-1.0	-1.9
-0.5	0.6
-0.1	1.7
1.2	3.3
3.1	1.6
1.3	1.6
2.0	0.8



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Pick K centers (randomly):

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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

```
> centers <- rbind(c(-1,-1),c(0,0))
```

> dist1 <- apply(x,1,function(x) sqrt(sum((x-centers[1,])^2</pre>

> dist2 <- apply(x,1,function(x) sqrt(sum((x-centers[2,])^2</pre>

Choose mean with smaller distance:

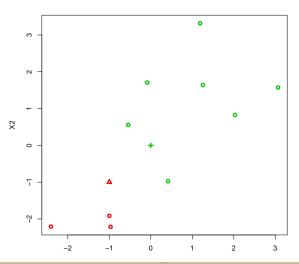
<i>X</i> ₁	<i>X</i> ₂	(-1,-1)	(0,0)
0.4	-1.0	1.4	1.1
-1.0	-2.2	1.2	2.4
-2.4	-2.2	1.9	3.3
-1.0	-1.9	0.9	2.2
-0.5	0.6	1.6	8.0
-0.1	1.7	2.9	1.7
1.2	3.3	4.8	3.5
3.1	1.6	4.8	3.4
1.3	1.6	3.5	2.1
2.0	0.8	3.5	2.2

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> dists <- cbind(dist1, dist2)</pre>

> cluster.ind <- apply(dists,1,which.min)</pre>

New clusters:

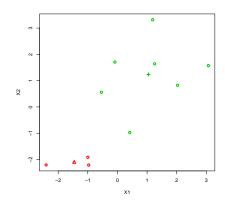


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Refit means for each cluster:

- 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

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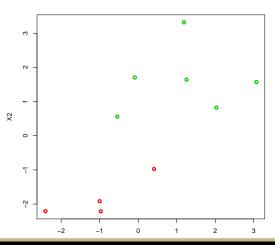
Choose mean with smaller distance:

<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

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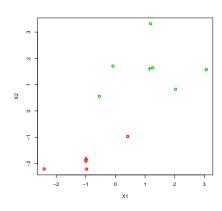
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New clusters:



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)



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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

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Select smallest distance and compare these clusters with previous:

Table: New Clusters

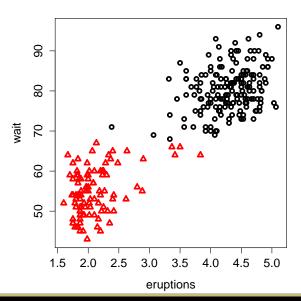
<i>X</i> ₁	<i>X</i> ₂	(-1.0, -1.8)	(1.2, 1.6)
0.4	-1.0	1.6	2.7
-1.0	-2.2	0.4	4.4
-2.4	-2.2	1.5	5.2
-1.0	-1.9	0.1	4.1
-0.5	0.6	2.4	2.0
-0.1	1.7	3.6	1.2
1.2	3.3	5.6	1.7
3.1	1.6	5.3	1.9
1.3	1.6	4.1	0.1
2.0	0.8	4.0	1.2

Table: 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

R has a function for K-means in the stats package; this is probably already loaded

- let's use this for the Old Faithful data
- > library(datasets)
- > faith.2 <- kmeans(faithful,2)</pre>
- > names(faith.2)
- > plot(faithful[,1],faithful[,2],col=faith.2\$clu
 - + pch=faith.2\$cluster, lwd=3)



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K-means can be used for *image* segmentation

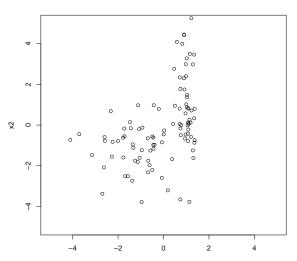
- partition image into multiple segments
- find boundaries of objects
- make art





K-Means Clustering

What is our data look like this?

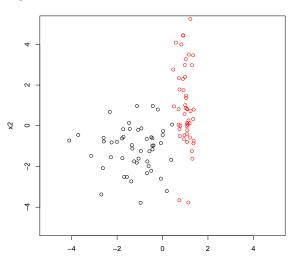


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K-Means Clustering

True clustering:



K-Means Clustering

K-means clustering (K = 2):

