
Expectation Maximization

Mixture Models

Clustering

Class 14. 6 Oct 2010

Administrivia

- ❑ Many homeworks still due
- ❑ Is everyone on the “projects” page?
 - Where are your project proposals?

Covered

- Learning distributions from data
 - Given a collection of examples from some data, estimate its distribution
 - Solution: Assign a model to the distribution
 - Learn parameters of model from data
- Complex models: Learning must be done using Expectation Maximization
- Following slides: An intuitive explanation using a simple example of multinomials

A Thought Experiment



COVERED

6 3 1 5 4 1 2 4 ...

- A person shoots a loaded dice repeatedly
- You observe the series of outcomes
- **You can form a good idea of how the dice is loaded**
 - Figure out what the probabilities of the various numbers are for dice
- $P(\text{number}) = \text{count}(\text{number}) / \text{sum}(\text{rolls})$
- This is a *maximum likelihood* estimate
 - Estimate that makes the observed sequence of numbers most probable

The Multinomial Distribution

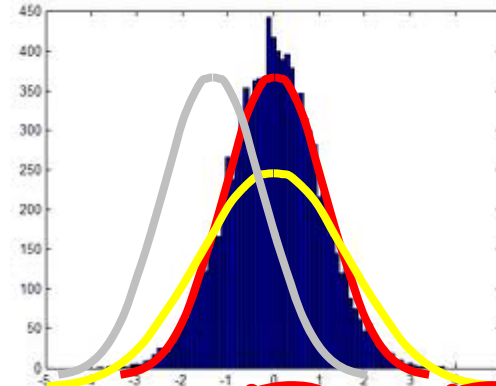
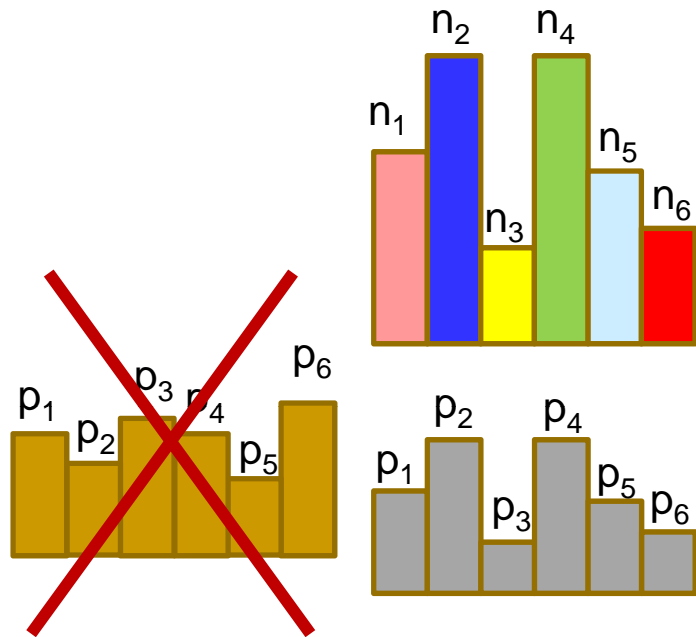
- A probability distribution over a discrete collection of items is a *Multinomial*

$$P(X : X \text{ belongs to a discrete set}) = P(X)$$

- E.g. the roll of dice
 - $X : X \text{ in } (1,2,3,4,5,6)$
- Or the toss of a coin
 - $X : X \text{ in } (\text{head}, \text{tails})$

COVERED

Maximum Likelihood Estimation



COVERED

- Basic principle: Assign a form to the distribution
 - E.g. a multinomial
 - Or a Gaussian
- Find the *distribution* that best fits the histogram of the data

Defining “Best Fit”

- The data are generated by draws from the distribution
 - I.e. the generating process draws from the distribution
- Assumption: The distribution has a high probability of generating the observed data
 - Not necessarily true
- Select the distribution that has the *highest* probability of generating the data
 - Should assign lower probability to less frequent observations and vice versa

COVERED

Maximum Likelihood Estimation: Multinomial

- Probability of generating $(n_1, n_2, n_3, n_4, n_5, n_6)$

$$P(n_1, n_2, n_3, n_4, n_5, n_6) = \text{Const} \prod_i p_i^{n_i}$$

- Find $p_1, p_2, p_3, p_4, p_5, p_6$ so that the above is maximized

- Alternately maximize

COVERED

$$\log(P(n_1, n_2, n_3, n_4, n_5, n_6)) = \log(\text{Const}) + \sum_i n_i \log(p_i)$$

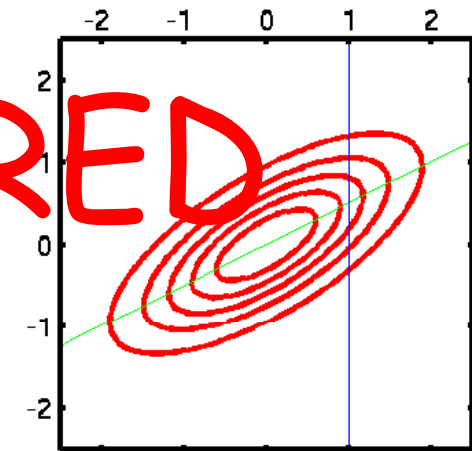
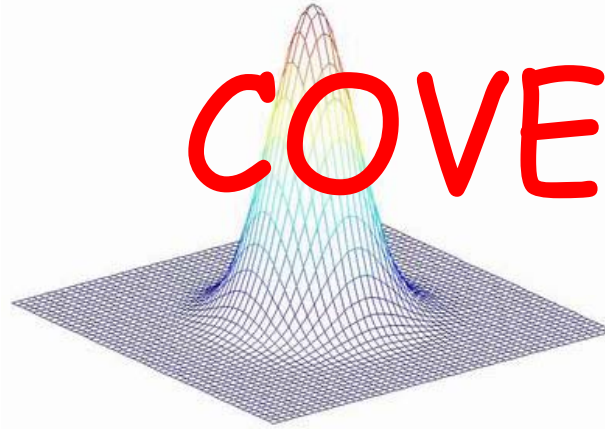
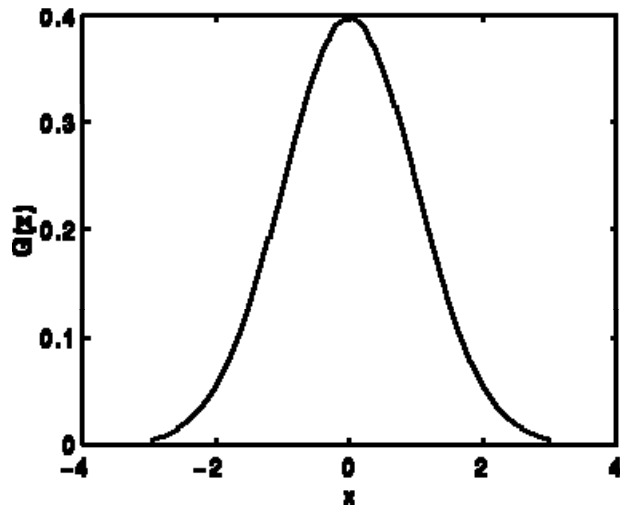
- Log() is a monotonic function
 - $\operatorname{argmax}_x f(x) = \operatorname{argmax}_x \log(f(x))$

- Solving for the probabilities gives us
 - Requires constrained optimization to ensure probabilities sum to 1

$$p_i = \frac{n_i}{\sum_j n_j}$$

**EVENTUALLY
ITS JUST
COUNTING!**

Segue: Gaussians



$$P(X) = N(X; \mu, \Theta) = \frac{1}{\sqrt{(2\pi)^d |\Theta|}} \exp(-0.5(X - \mu)^T \Theta^{-1} (X - \mu))$$

- Parameters of a Gaussian:
 - Mean μ , Covariance Θ

Maximum Likelihood: Gaussian

- Given a collection of observations (X_1, X_2, \dots) , estimate mean μ and covariance Θ

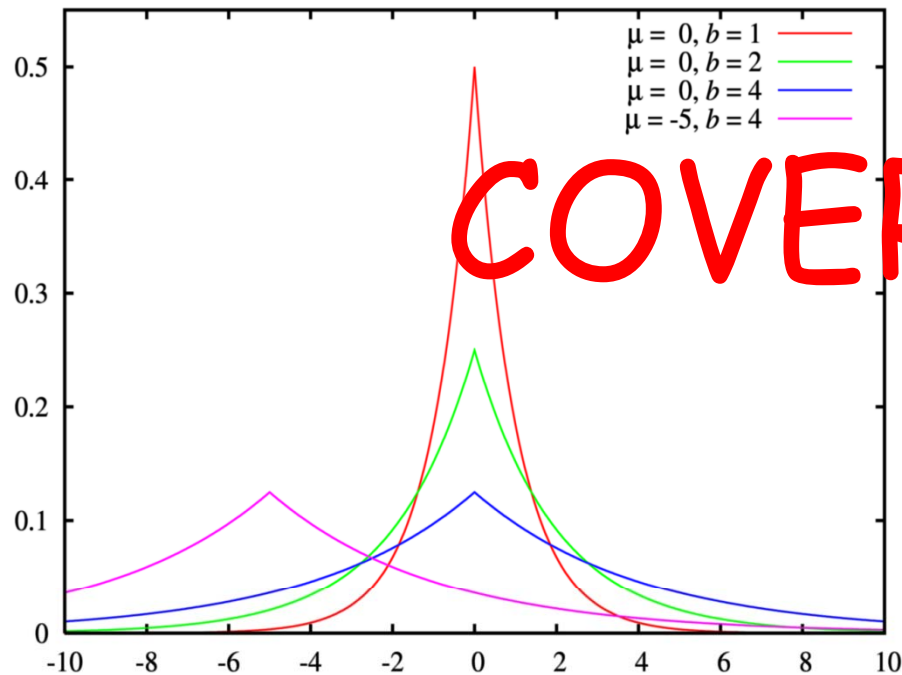
$$P(X_1, X_2, \dots) = \prod_i \frac{1}{\sqrt{(2\pi)^d |\Theta|}} \exp(-0.5(X_i - \mu)^T \Theta^{-1} (X_i - \mu))$$
$$\log(P(X_1, X_2, \dots)) = C - 0.5 \sum_i (\log(|\Theta|) + (X_i - \mu)^T \Theta^{-1} (X_i - \mu))$$

- Maximizing w.r.t μ and Θ gives us

$$\mu = \frac{1}{N} \sum_i X_i \quad \Theta = \frac{1}{N} \sum_i (X_i - \mu)(X_i - \mu)^T$$

ITS STILL
JUST
COUNTING!

Laplacian



$$P(x) = L(x; \mu, b) = \frac{1}{2b} \exp\left(-\frac{|x - \mu|}{b}\right)$$

- Parameters: Mean μ , scale b ($b > 0$)

Maximum Likelihood: Laplacian

- Given a collection of observations (x_1, x_2, \dots) , estimate mean μ and scale b

$$\log(P(x_1, x_2, \dots)) = C - N \log(b) - \sum_i \frac{|x_i - \mu|}{b}$$

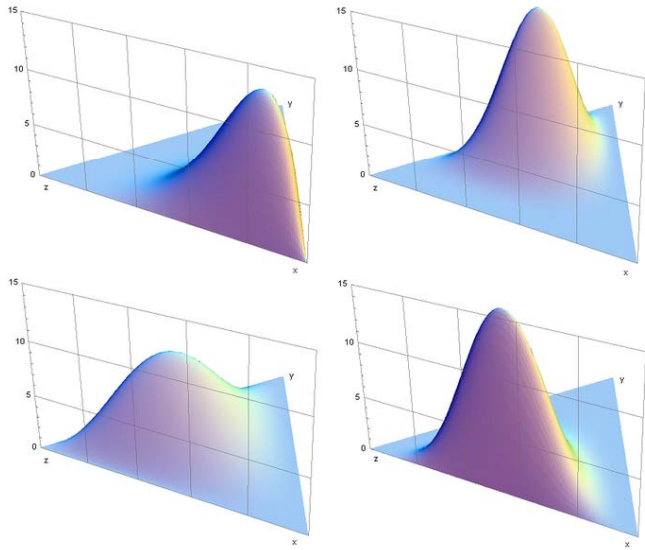
COVERED

- Maximizing w.r.t μ and b gives us

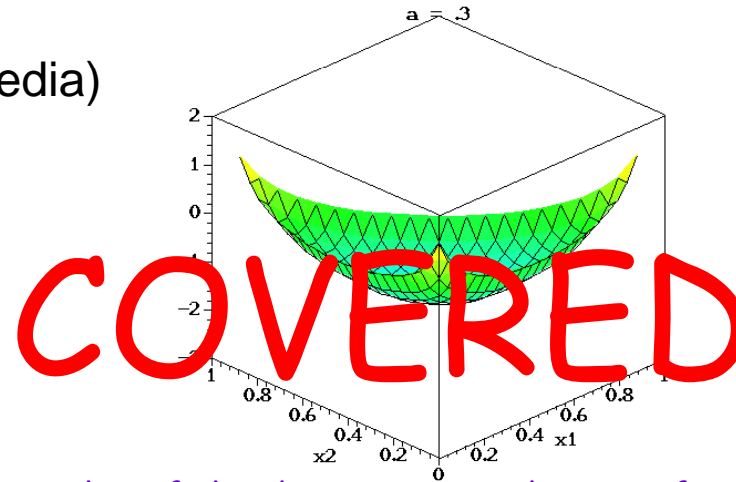
$$\mu = \frac{1}{N} \sum_i x_i \qquad b = \frac{1}{N} \sum_i |x_i - \mu|$$

Dirichlet

(from wikipedia)



$K=3$. Clockwise from top left:
 $\alpha=(6, 2, 2), (3, 7, 5), (6, 2, 6), (2, 3, 4)$



log of the density as we change a from $\alpha=(0.3, 0.3, 0.3)$ to $(2.0, 2.0, 2.0)$, keeping all the individual a_i 's equal to each other.

- Parameters are α s
 - Determine mode and curvature
- Defined only of probability vectors
 - $X = [x_1 \ x_2 \ .. \ x_K], \sum_i x_i = 1, x_i \geq 0$ for all i

$$P(X) = D(X; \alpha) = \frac{\prod \Gamma(\alpha_i)}{\Gamma\left(\sum_i \alpha_i\right)} \prod_i x_i^{\alpha_i - 1}$$

Maximum Likelihood: Dirichlet

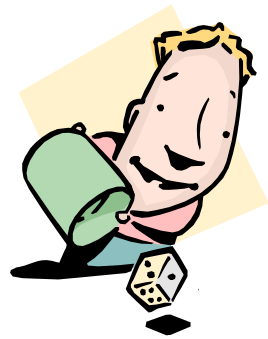
- Given a collection of observations (X_1, X_2, \dots) , estimate α

COVERED

$$\log(P(X_1, X_2, \dots)) = \sum_j \sum_i (\alpha_i - 1) \log(X_{j,i}) + N \sum_i \log(\Gamma(\alpha_i)) - N \log\left(\Gamma\left(\sum_i \alpha_i\right)\right)$$

- No closed form solution for α s.
 - Needs gradient ascent
- Several distributions have this property: the ML estimate of their parameters have no closed form solution

Continuing the Thought Experiment



6 3 1 5 4 1 2 4 ...

COVERED



4 4 1 6 3 2 1 2 ...

- Two persons shoot loaded dice repeatedly
 - The dice are differently loaded for the two of them
- We observe the series of outcomes for both persons
- **How to determine the probability distributions of the two dice?**

Estimating Probabilities

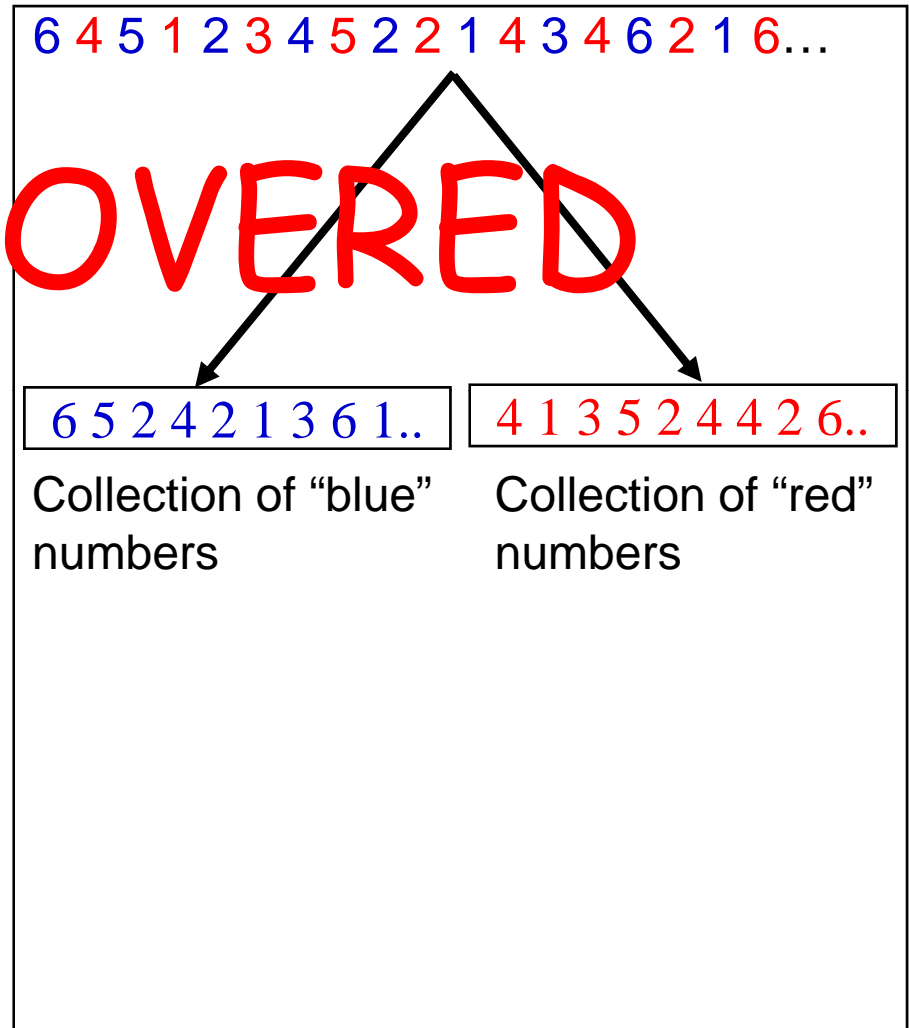
- Observation: The sequence of numbers from the two dice
 - As indicated by the colors, we know who rolled what number

6 4 5 1 2 3 4 5 2 2 1 4 3 4 6 2 1 6...

COVERED

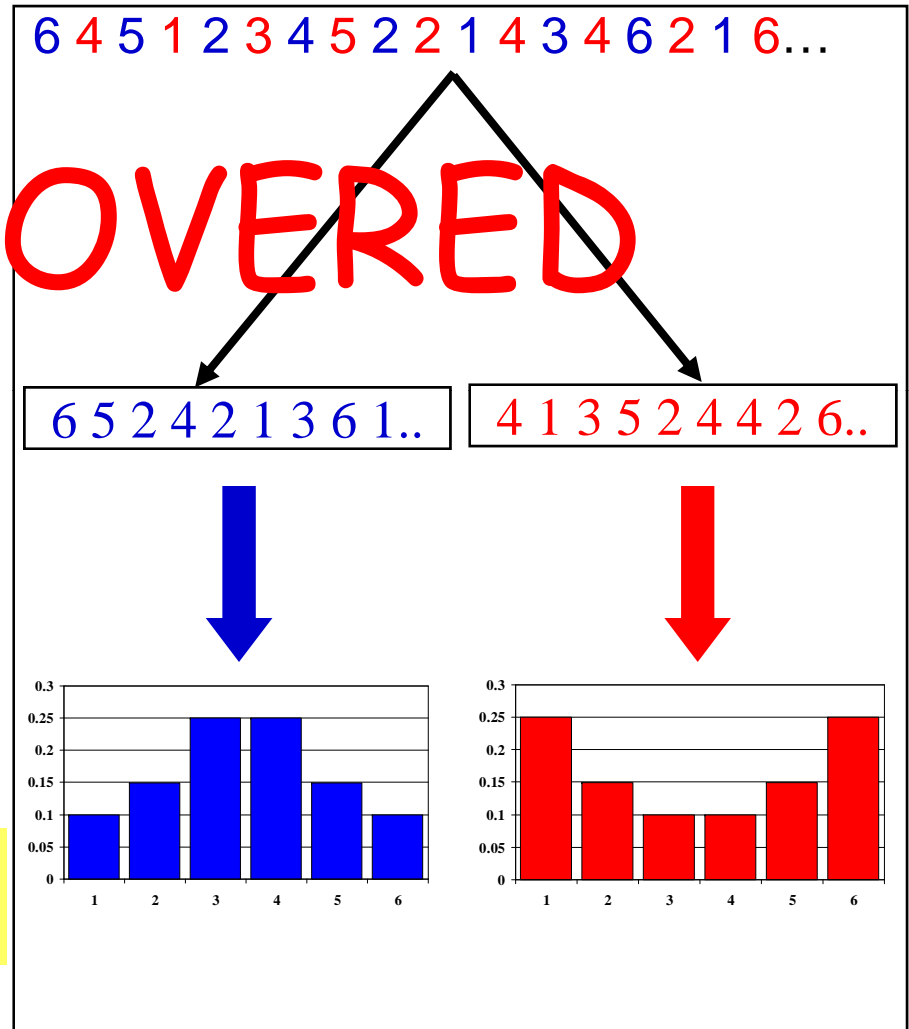
Estimating Probabilities

- Observation: The sequence of numbers from the two dice
 - As indicated by the colors, we know who rolled what number
- Segregation: Separate the blue observations from the red



Estimating Probabilities

- Observation: The sequence of numbers from the two dice
 - As indicated by the colors, we know who rolled what number
- Segregation: Separate the blue observations from the red
- From each set compute probabilities for each of the 6 possible outcomes



$$P(\text{number}) = \frac{\text{no. of times number was rolled}}{\text{total number of observed rolls}}$$

A Thought Experiment

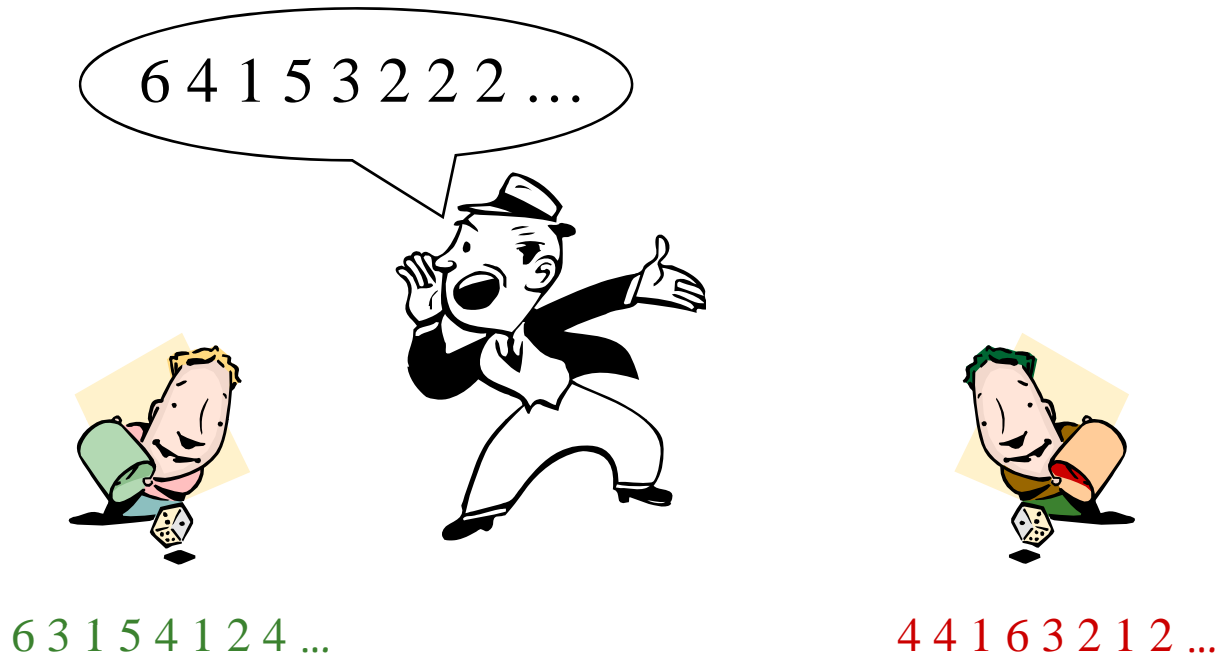


6 3 1 5 4 1 2 4 ...

4 4 1 6 3 2 1 2 ...

- Now imagine that you cannot observe the dice yourself
- Instead there is a “caller” who randomly calls out the outcomes
 - 40% of the time he calls out the number from the left shooter, and 60% of the time, the one from the right (and you know this)
- At any time, you do not know which of the two he is calling out
- How do you determine the probability distributions for the two dice?

A Thought Experiment



- How do you now determine the probability distributions for the two sets of dice ...
- .. If you do not even know what fraction of time the blue numbers are called, and what fraction are red?

A Mixture Multinomial

- The caller will call out a number X in any given callout IF
 - He selects “RED”, and the Red die rolls the number X
 - OR
 - He selects “BLUE” and the Blue die rolls the number X
- $P(X) = P(\text{Red})P(X|\text{Red}) + P(\text{Blue})P(X|\text{Blue})$
 - E.g. $P(6) = P(\text{Red})P(6|\text{Red}) + P(\text{Blue})P(6|\text{Blue})$
- A distribution that *combines* (or *mixes*) multiple multinomials is a *mixture* multinomial

$$P(X) = \sum_Z P(Z)P(X|Z)$$

Mixture weights

Component multinomials

Mixture Distributions

$$P(X) = \sum_Z P(Z)P(X | Z)$$

Mixture weights Component distributions

Mixture Gaussian

$$P(X) = \sum_Z P(Z)N(X; \mu_z, \Theta_z)$$

Mixture of Gaussians and Laplacians

$$P(X) = \sum_Z P(Z)N(X; \mu_z, \Theta_z) + \sum_Z P(Z) \prod_i L(X_i; \mu_z, b_{z,i})$$

- Mixture distributions mix several component distributions
 - Component distributions may be of varied type
- Mixing weights must sum to 1.0
- Component distributions integrate to 1.0
- Mixture distribution integrates to 1.0

Maximum Likelihood Estimation

- For our problem:
$$P(X) = \sum_Z P(Z)P(X | Z)$$
 - $Z = \text{color of dice}$

$$P(n_1, n_2, n_3, n_4, n_5, n_6) = \text{Const} \prod_X P(X)^{n_x} = \text{Const} \prod_X \left(\sum_Z P(Z)P(X | Z) \right)^{n_x}$$

- Maximum likelihood solution: Maximize

$$\log(P(n_1, n_2, n_3, n_4, n_5, n_6)) = \log(\text{Const}) + \sum_X n_x \log \left(\sum_Z P(Z)P(X | Z) \right)$$

- No closed form solution (summation inside log)!
 - In general ML estimates for mixtures do not have a closed form
 - USE EM!

Expectation Maximization

- It is possible to estimate all parameters in this setup using the Expectation Maximization (or EM) algorithm
- First described in a landmark paper by Dempster, Laird and Rubin
 - Maximum Likelihood Estimation from incomplete data, via the EM Algorithm, Journal of the Royal Statistical Society, Series B, 1977
- Much work on the algorithm since then
- The principles behind the algorithm existed for several years prior to the landmark paper, however.

Expectation Maximization

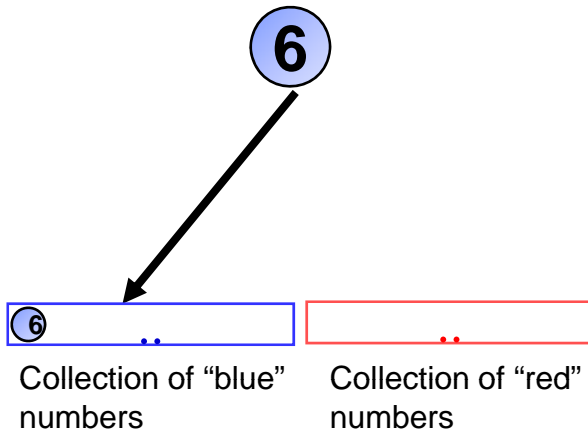
- Iterative solution
- Get some initial estimates for all parameters
 - Dice shooter example: This includes probability distributions for dice AND the probability with which the caller selects the dice
- Two steps that are iterated:
 - **Expectation Step:** Estimate statistically, the values of *unseen* variables
 - **Maximization Step:** Using the estimated values of the unseen variables as truth, estimates of the model parameters

EM: The auxiliary function

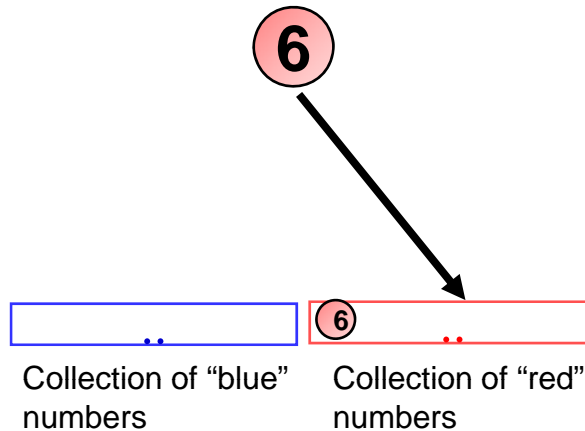
- EM iteratively optimizes the following auxiliary function
- $Q(\theta, \theta') = \sum_Z P(Z|X, \theta') \log(P(Z, X | \theta))$
 - Z are the unseen variables
 - Assuming Z is discrete (may not be)
- θ' are the parameter estimates from the previous iteration
- θ are the estimates to be obtained in the current iteration

Expectation Maximization as counting

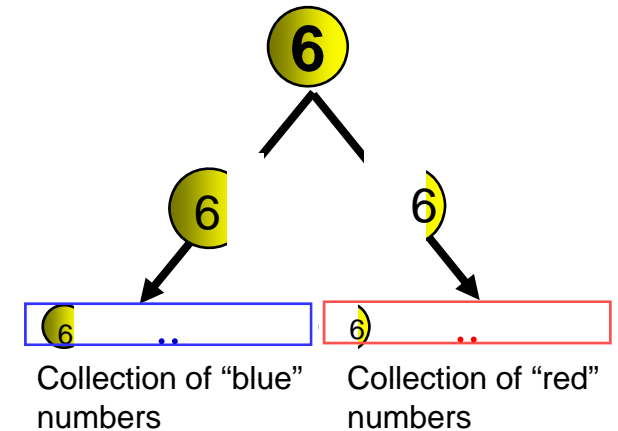
Instance from blue dice



Instance from red dice



Dice unknown



- Hidden variable: Z
 - Dice: The identity of the dice whose number has been called out
- If we knew Z for every observation, we could estimate all terms
 - By adding the observation to the correct bin
- Unfortunately, we do not know Z – it is hidden from us!
- Solution: FRAGMENT THE OBSERVATION

Fragmenting the Observation

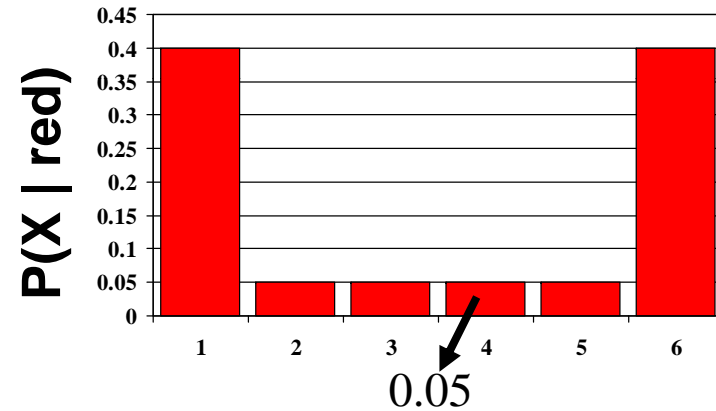
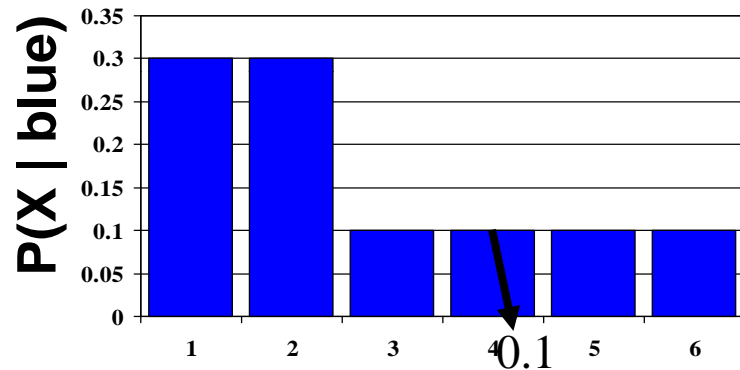
- EM is an iterative algorithm
 - At each time there is a *current* estimate of parameters
- The “size” of the fragments is proportional to the *a posteriori probability* of the component distributions
 - The *a posteriori* probabilities of the various values of Z are computed using Bayes' rule:

$$P(Z | X) = \frac{P(X | Z)P(Z)}{P(X)} = CP(X | Z)P(Z)$$

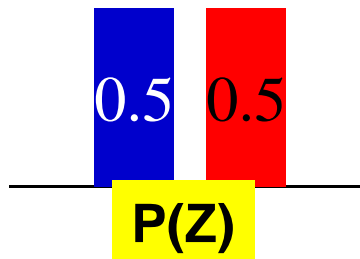
- Every dice gets a fragment of size $P(\text{dice} | \text{number})$

Expectation Maximization

- Hypothetical Dice Shooter Example:
- We obtain an initial estimate for the probability distribution of the two sets of dice (somehow):



- We obtain an initial estimate for the probability with which the caller calls out the two shooters (somehow)



Expectation Maximization

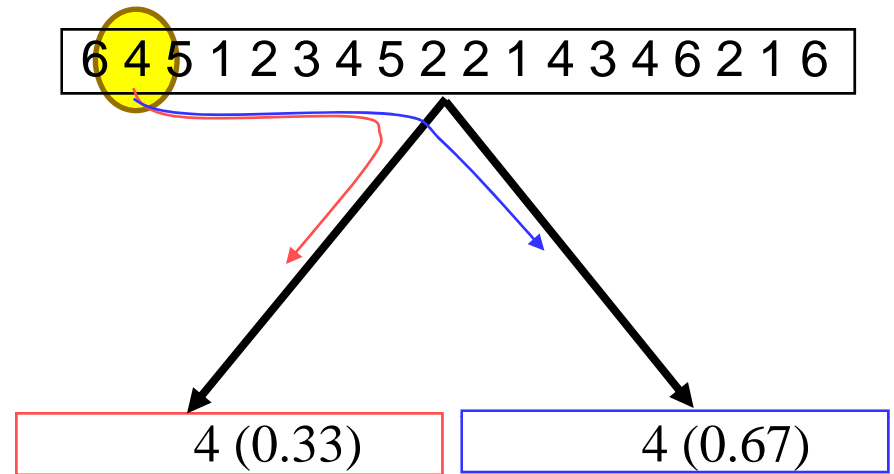
- Hypothetical Dice Shooter Example:
- Initial estimate:
 - $P(\text{blue}) = P(\text{red}) = 0.5$
 - $P(4 \mid \text{blue}) = 0.1$, for $P(4 \mid \text{red}) = 0.05$
- Caller has just called out 4
- Posterior probability of colors:

$$P(\text{red} \mid X = 4) = CP(X = 4 \mid Z = \text{red})P(Z = \text{red}) = C \times 0.05 \times 0.5 = C0.025$$

$$P(\text{blue} \mid X = 4) = CP(X = 4 \mid Z = \text{blue})P(Z = \text{blue}) = C \times 0.1 \times 0.5 = C0.05$$

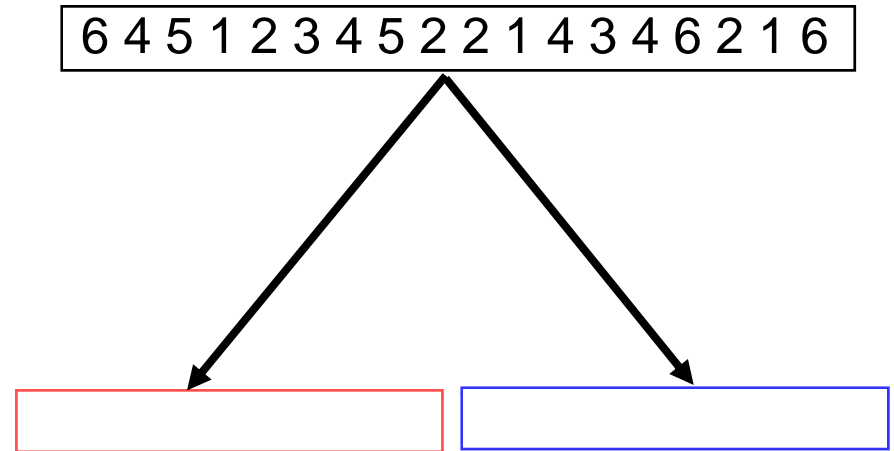
$$\text{Normalizing : } P(\text{red} \mid X = 4) = 0.33; \quad P(\text{blue} \mid X = 4) = 0.67$$

Expectation **Maximization**



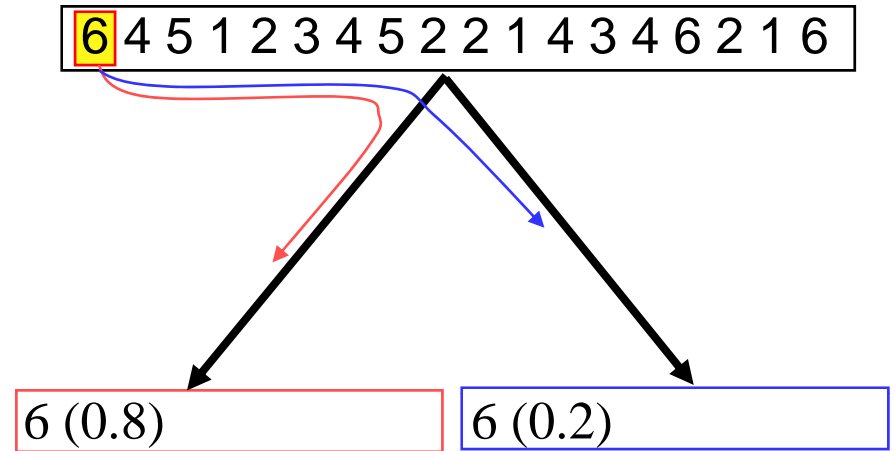
Expectation **Maximization**

- Every observed roll of the dice contributes to both “Red” and “Blue”



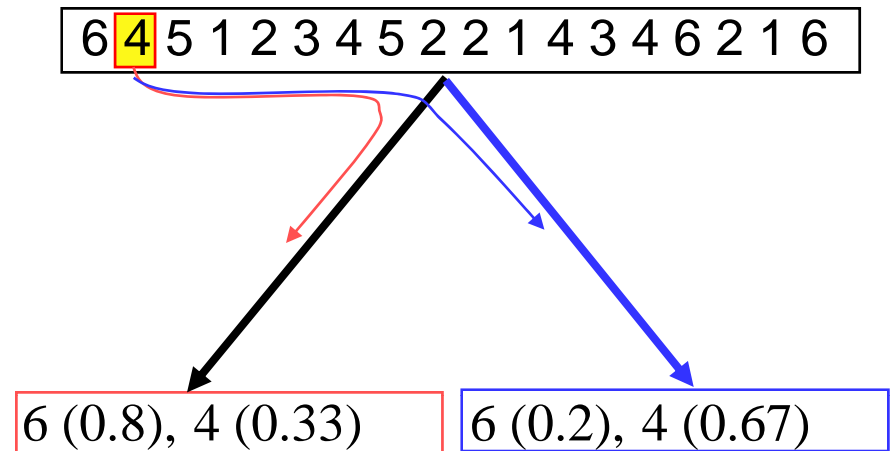
Expectation **Maximization**

- Every observed roll of the dice contributes to both “Red” and “Blue”



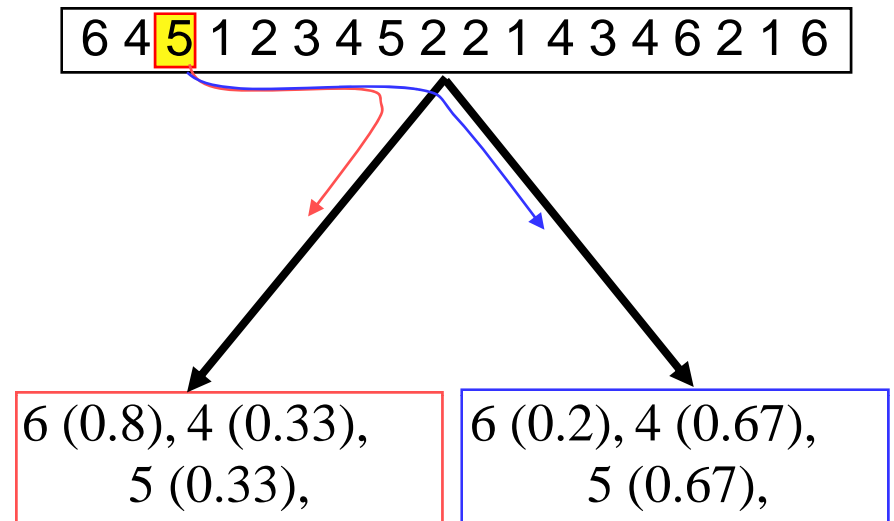
Expectation **Maximization**

- Every observed roll of the dice contributes to both “Red” and “Blue”



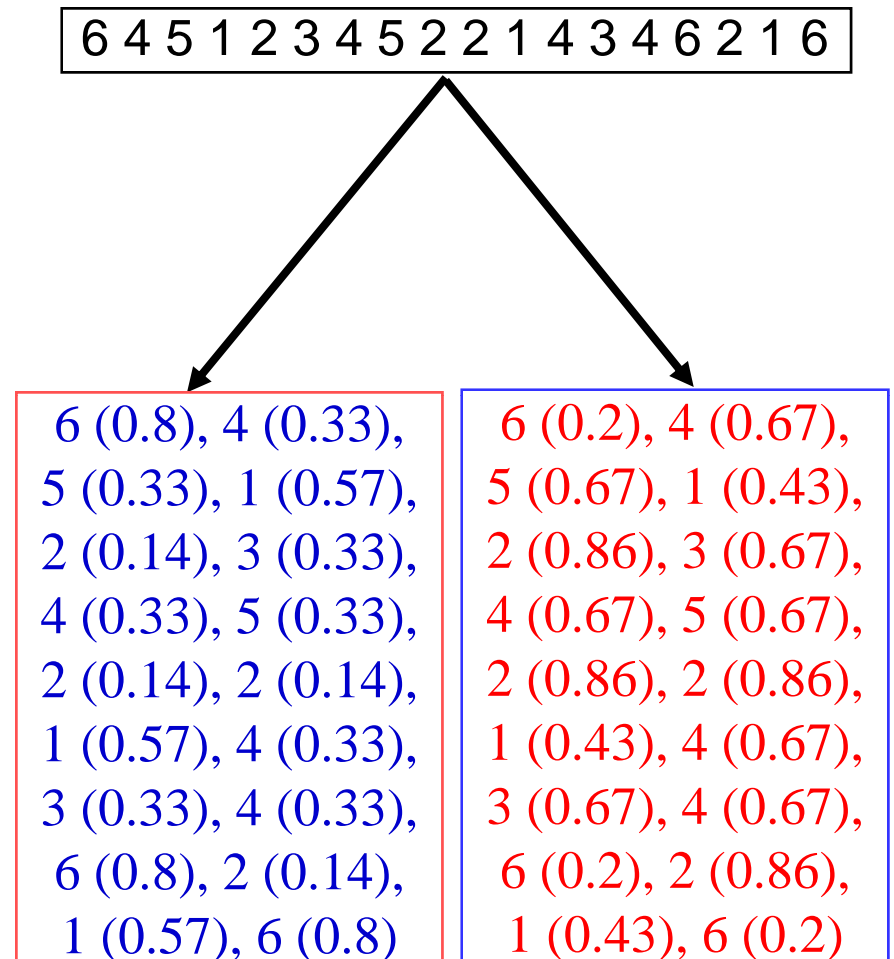
Expectation **Maximization**

- Every observed roll of the dice contributes to both “Red” and “Blue”



Expectation **Maximization**

- Every observed roll of the dice contributes to both “Red” and “Blue”



Expectation **Maximization**

- Every observed roll of the dice contributes to both “Red” and “Blue”
- Total count for “Red” is the sum of all the posterior probabilities in the red column
 - 7.31
- Total count for “Blue” is the sum of all the posterior probabilities in the blue column
 - 10.69
 - Note: $10.69 + 7.31 = 18 =$ the total number of instances

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69³⁷

Expectation **Maximization**

- Total count for “Red” : 7.31
- Red:
 - Total count for 1: 1.71

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₃₈

Expectation **Maximization**

- Total count for “Red” : 7.31
- Red:
 - Total count for 1: 1.71
 - Total count for 2: 0.56

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69³⁹

Expectation **Maximization**

- Total count for “Red” : 7.31
- Red:
 - Total count for 1: 1.71
 - Total count for 2: 0.56
 - Total count for 3: 0.66

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₀

Expectation **Maximization**

- Total count for “Red” : 7.31
- Red:
 - Total count for 1: 1.71
 - Total count for 2: 0.56
 - Total count for 3: 0.66
 - Total count for 4: 1.32

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₁

Expectation **Maximization**

- Total count for “Red” : 7.31
- Red:
 - Total count for 1: 1.71
 - Total count for 2: 0.56
 - Total count for 3: 0.66
 - Total count for 4: 1.32
 - Total count for 5: 0.66

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₂

Expectation **Maximization**

- Total count for “Red” : 7.31
- Red:
 - Total count for 1: 1.71
 - Total count for 2: 0.56
 - Total count for 3: 0.66
 - Total count for 4: 1.32
 - Total count for 5: 0.66
 - Total count for 6: 2.4

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₃

Expectation **Maximization**

- Total count for “Red” : 7.31
- Red:
 - Total count for 1: 1.71
 - Total count for 2: 0.56
 - Total count for 3: 0.66
 - Total count for 4: 1.32
 - Total count for 5: 0.66
 - Total count for 6: 2.4
- **Updated probability of Red dice:**
 - $P(1 | \text{Red}) = 1.71/7.31 = 0.234$
 - $P(2 | \text{Red}) = 0.56/7.31 = 0.077$
 - $P(3 | \text{Red}) = 0.66/7.31 = 0.090$
 - $P(4 | \text{Red}) = 1.32/7.31 = 0.181$
 - $P(5 | \text{Red}) = 0.66/7.31 = 0.090$
 - $P(6 | \text{Red}) = 2.40/7.31 = 0.328$

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₄

Expectation **Maximization**

- Total count for “Blue” : 10.69
- Blue:
 - Total count for 1: 1.29

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₅

Expectation **Maximization**

- Total count for “Blue” : 10.69
- Blue:
 - Total count for 1: 1.29
 - Total count for 2: 3.44

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₆

Expectation **Maximization**

- Total count for “Blue” : 10.69
- Blue:
 - Total count for 1: 1.29
 - Total count for 2: 3.44
 - Total count for 3: 1.34

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₇

Expectation **Maximization**

- Total count for “Blue” : 10.69
- Blue:
 - Total count for 1: 1.29
 - Total count for 2: 3.44
 - Total count for 3: 1.34
 - Total count for 4: 2.68

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₈

Expectation **Maximization**

- Total count for “Blue” : 10.69
- Blue:
 - Total count for 1: 1.29
 - Total count for 2: 3.44
 - Total count for 3: 1.34
 - Total count for 4: 2.68
 - Total count for 5: 1.34

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₄₉

Expectation **Maximization**

- Total count for “Blue” : 10.69
- Blue:
 - Total count for 1: 1.29
 - Total count for 2: 3.44
 - Total count for 3: 1.34
 - Total count for 4: 2.68
 - Total count for 5: 1.34
 - Total count for 6: 0.6

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₅₀

Expectation **Maximization**

- Total count for “Blue” : 10.69
- Blue:
 - Total count for 1: 1.29
 - Total count for 2: 3.44
 - Total count for 3: 1.34
 - Total count for 4: 2.68
 - Total count for 5: 1.34
 - Total count for 6: 0.6
- **Updated probability of Blue dice:**
 - $P(1 | \text{Blue}) = 1.29/11.69 = 0.122$
 - $P(2 | \text{Blue}) = 0.56/11.69 = 0.322$
 - $P(3 | \text{Blue}) = 0.66/11.69 = 0.125$
 - $P(4 | \text{Blue}) = 1.32/11.69 = 0.250$
 - $P(5 | \text{Blue}) = 0.66/11.69 = 0.125$
 - $P(6 | \text{Blue}) = 2.40/11.69 = 0.056$

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₅₁

Expectation **Maximization**

- Total count for “Red” : 7.31
- Total count for “Blue” : 10.69
- Total instances = 18
 - Note $7.31 + 10.69 = 18$
- We also revise our estimate for the probability that the caller calls out Red or Blue
 - i.e the fraction of times that he calls Red and the fraction of times he calls Blue
- $P(Z=\text{Red}) = 7.31/18 = 0.41$
- $P(Z=\text{Blue}) = 10.69/18 = 0.59$

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

7.31

10.69₅₂

The updated values

- Probability of Red dice:

- $P(1 | \text{Red}) = 1.71/7.31 = 0.234$
- $P(2 | \text{Red}) = 0.56/7.31 = 0.077$
- $P(3 | \text{Red}) = 0.66/7.31 = 0.090$
- $P(4 | \text{Red}) = 1.32/7.31 = 0.181$
- $P(5 | \text{Red}) = 0.66/7.31 = 0.090$
- $P(6 | \text{Red}) = 2.40/7.31 = 0.328$

- Probability of Blue dice:

- $P(1 | \text{Blue}) = 1.29/11.69 = 0.122$
- $P(2 | \text{Blue}) = 0.56/11.69 = 0.322$
- $P(3 | \text{Blue}) = 0.66/11.69 = 0.125$
- $P(4 | \text{Blue}) = 1.32/11.69 = 0.250$
- $P(5 | \text{Blue}) = 0.66/11.69 = 0.125$
- $P(6 | \text{Blue}) = 2.40/11.69 = 0.056$

- $P(Z=\text{Red}) = 7.31/18 = 0.41$

- $P(Z=\text{Blue}) = 10.69/18 = 0.59$

Called	P(red X)	P(blue X)
6	.8	.2
4	.33	.67
5	.33	.67
1	.57	.43
2	.14	.86
3	.33	.67
4	.33	.67
5	.33	.67
2	.14	.86
2	.14	.86
1	.57	.43
4	.33	.67
3	.33	.67
4	.33	.67
6	.8	.2
2	.14	.86
1	.57	.43
6	.8	.2

THE UPDATED VALUES CAN BE USED TO REPEAT THE PROCESS. ESTIMATION IS AN ITERATIVE PROCESS

The Dice Shooter Example



6 3 1 5 4 1 2 4 ...

4 4 1 6 3 2 1 2 ...

1. Initialize $P(Z)$, $P(X | Z)$
2. Estimate $P(Z | X)$ for each Z , for each called out number
 - Assign X with to value of Z , with weight $P(Z | X)$
3. Re-estimate $P(X | Z)$ for every value of X and Z
4. Re-estimate $P(Z)$
5. If not converged, return to 2

In Squiggles

- Given a sequence of observations O_1, O_2, \dots
 - N_x is the number of observations of number X
- Initialize $P(Z), P(X|Z)$ for dice Z and numbers X
- Iterate:

- For each number X :

$$P(Z | X) = \frac{P(X | Z)P(Z)}{\sum_{Z'} P(Z')P(X | Z')}$$

- Update:

$$P(X | Z) = \frac{\sum_{O \text{ such that } O=X} P(Z | O)}{\sum_O P(Z | O)} = \frac{N_X P(Z | X)}{\sum_X N_X P(Z | X)}$$

$$P(Z) = \frac{\sum_X N_X P(Z | X)}{\sum_{Z'} \sum_X N_X P(Z' | X)}$$

Solutions may not be unique

- The EM algorithm will give us one of many solutions, all equally valid!

- The probability of 6 being called out:

$$P(6) = \alpha P(6 | red) + \beta P(6 | blue) = \alpha P_r + \beta P_b$$

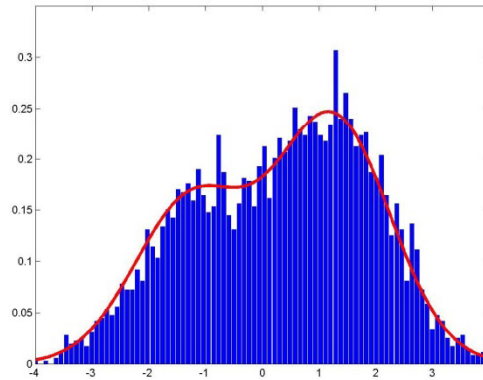
- Assigns P_r as the probability of 6 for the red die
- Assigns P_b as the probability of 6 for the blue die

- The following too is a valid solution

$$P(6) = 1.0(\alpha P_r + \beta P_b) + 0.0 \text{ anything}$$

- Assigns 1.0 as the a priori probability of the red die
 - Assigns 0.0 as the probability of the blue die
- The solution is NOT unique

A More Complex Model



$$P(X) = \sum_k P(k) N(X; \mu_k, \Theta_k) = \sum_k \frac{P(k)}{\sqrt{(2\pi)^d |\Theta_k|}} \exp\left(-0.5(X - \mu_k)^T \Theta_k^{-1} (X - \mu_k)\right)$$

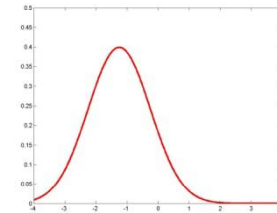
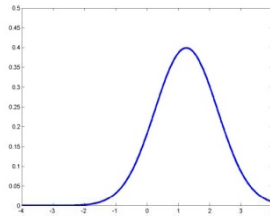
- Gaussian mixtures are often good models for the distribution of multivariate data
- Problem: Estimating the parameters, given a collection of data

Gaussian Mixtures: Generating model

$$P(X) = \sum_k P(k)N(X; \mu_k, \Theta_k)$$



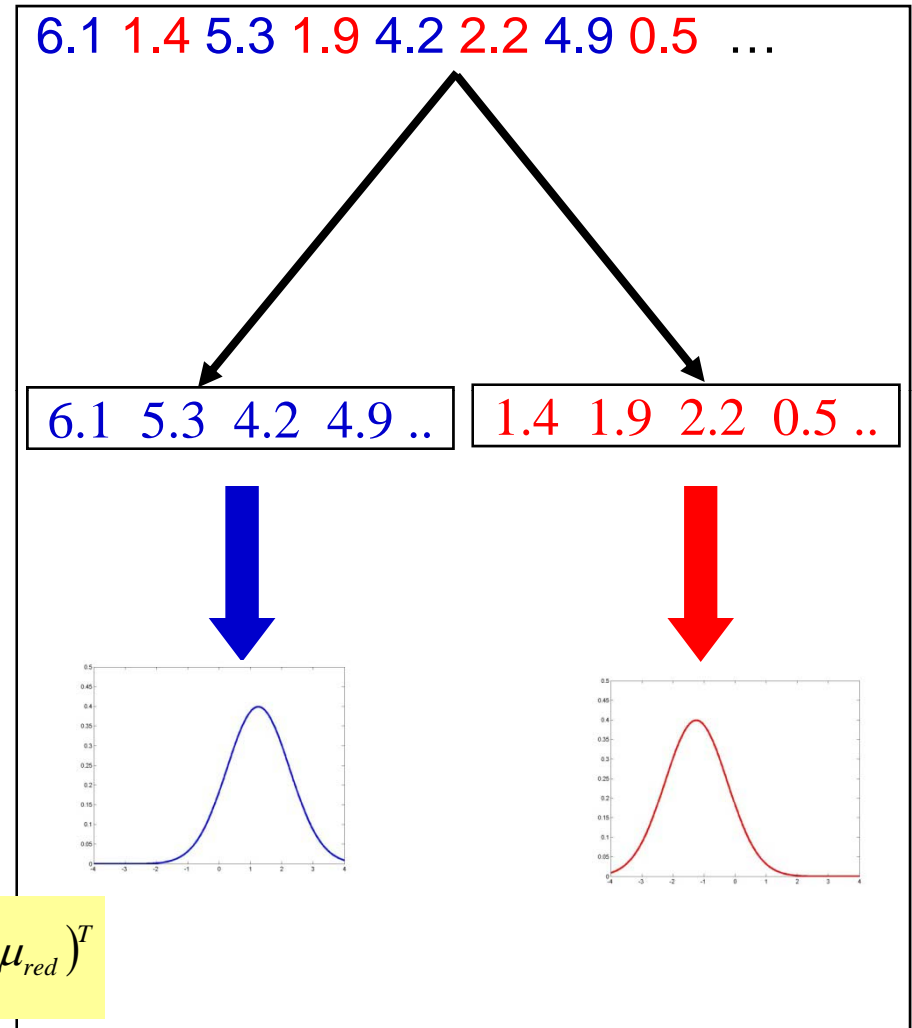
6.1 1.4 5.3 1.9 4.2 2.2 4.9 0.5



- The caller now has two Gaussians
 - At each draw he randomly selects a Gaussian, by the mixture weight distribution
 - He then draws an observation from that Gaussian
 - Much like the dice problem (only the outcomes are now real numbers and can be anything)

Estimating GMM with complete information

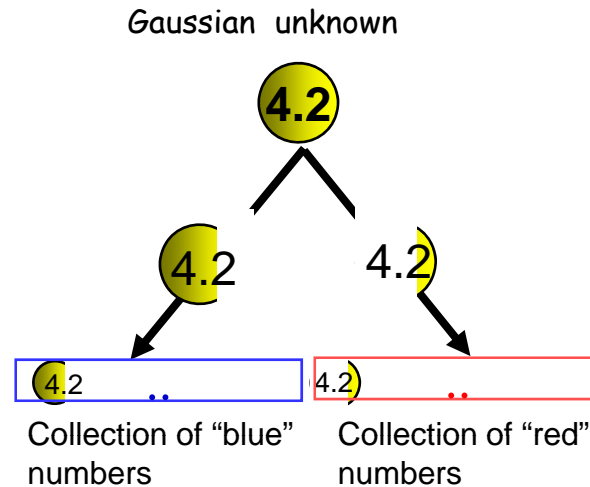
- Observation: A collection of numbers drawn from a mixture of 2 Gaussians
 - As indicated by the colors, we know which Gaussian generated what number
- Segregation: Separate the blue observations from the red
- From each set compute parameters for that Gaussian



$$\mu_{red} = \frac{1}{N_{red}} \sum_{i \in red} X_i \quad \Theta_{red} = \frac{1}{N_{red}} \sum_{i \in red} (X_i - \mu_{red})(X_i - \mu_{red})^T$$

$$P(red) = \frac{N_{red}}{N}$$

Fragmenting the observation



- The identity of the Gaussian is not known!
- Solution: **Fragment the observation**
- Fragment size proportional to *a posteriori* probability

$$P(k | X) = \frac{P(X | k)P(k)}{\sum_{k'} P(k')P(X | k')} = \frac{P(k)N(X; \mu_k, \Theta_k)}{\sum_{k'} P(k')N(X; \mu_{k'}, \Theta_{k'})}$$

Expectation **Maximization**

- Initialize $P(k)$, μ_k and Θ_k for both Gaussians
 - Important how we do this
 - Typical solution: Initialize means randomly, Θ_k as the global covariance of the data and $P(k)$ uniformly
- Compute fragment sizes for each Gaussian, for each observation

Number	P(red X)	P(blue X)
6.1	.81	.19
1.4	.33	.67
5.3	.75	.25
1.9	.41	.59
4.2	.64	.36
2.2	.43	.57
4.9	.66	.34
0.5	.05	.95

$$P(k | X) = \frac{P(k)N(X; \mu_k, \Theta_k)}{\sum_{k'} P(k')N(X; \mu_{k'}, \Theta_{k'})}$$

Expectation **Maximization**

- **Each observation contributes only as much as its fragment size to each statistic**

- $\text{Mean}(\text{red}) = (6.1 \cdot 0.81 + 1.4 \cdot 0.33 + 5.3 \cdot 0.75 + 1.9 \cdot 0.41 + 4.2 \cdot 0.64 + 2.2 \cdot 0.43 + 4.9 \cdot 0.66 + 0.5 \cdot 0.05) / (0.81 + 0.33 + 0.75 + 0.41 + 0.64 + 0.43 + 0.66 + 0.05) = 17.05 / 4.08 = 4.18$

Number	P(red X)	P(blue X)
6.1	.81	.19
1.4	.33	.67
5.3	.75	.25
1.9	.41	.59
4.2	.64	.36
2.2	.43	.57
4.9	.66	.34
0.5	.05	.95

4.08

3.92

- $\text{Var}(\text{red}) = ((6.1-4.18)^2 \cdot 0.81 + (1.4-4.18)^2 \cdot 0.33 + (5.3-4.18)^2 \cdot 0.75 + (1.9-4.18)^2 \cdot 0.41 + (4.2-4.18)^2 \cdot 0.64 + (2.2-4.18)^2 \cdot 0.43 + (4.9-4.18)^2 \cdot 0.66 + (0.5-4.18)^2 \cdot 0.05) / (0.81 + 0.33 + 0.75 + 0.41 + 0.64 + 0.43 + 0.66 + 0.05)$

$$P(\text{red}) = \frac{4.08}{8}$$

EM for Gaussian Mixtures

1. Initialize $P(k)$, μ_k and Θ_k for all Gaussians
2. For each observation X compute *a posteriori* probabilities for all Gaussian

$$P(k | X) = \frac{P(k)N(X; \mu_k, \Theta_k)}{\sum_{k'} P(k')N(X; \mu_{k'}, \Theta_{k'})}$$

3. Update mixture weights, means and variances for all Gaussians

$$P(k) = \frac{\sum_X P(k|X)}{N}$$

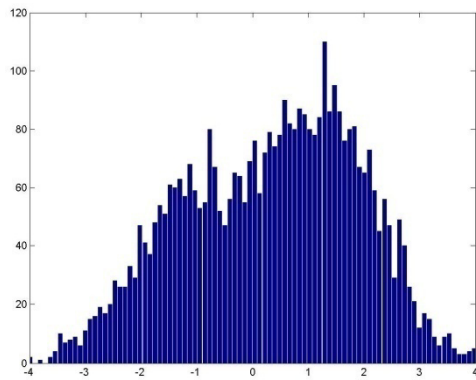
$$\mu_k = \frac{\sum_X P(k|X) X}{\sum_X P(k|X)}$$

$$\Theta_k = \frac{\sum_X P(k|X) (X - \mu_k)^2}{\sum_X P(k|X)}$$

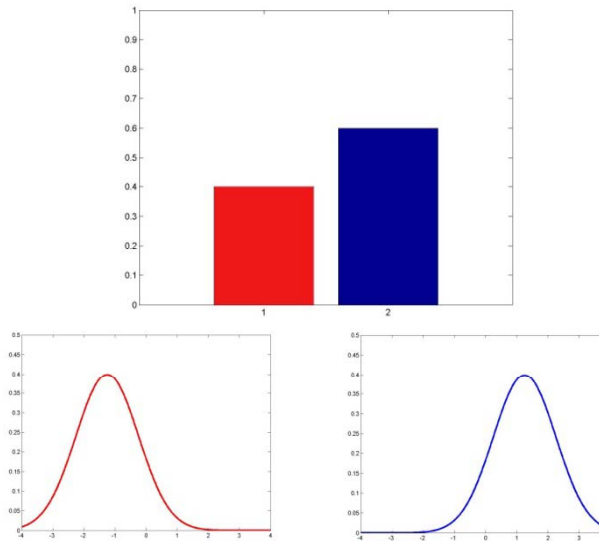
4. If not converged, return to 2

EM estimation of Gaussian Mixtures

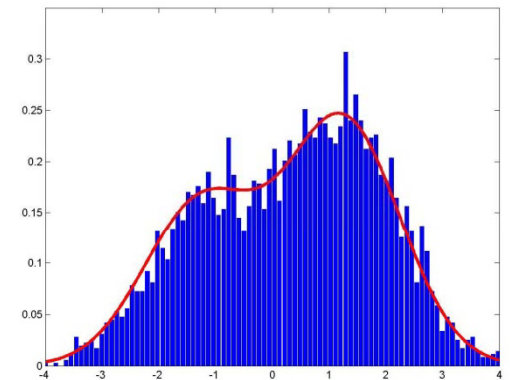
- An Example



Histogram of 4000 instances of a randomly generated data



Individual parameters of a two-Gaussian mixture estimated by EM



Two-Gaussian mixture estimated by EM

Expectation Maximization

- The same principle can be extended to mixtures of other distributions.
- E.g. Mixture of Laplacians: Laplacian parameters become

$$\mu_k = \frac{1}{\sum_x P(k|x)} \sum_x P(k|x)x \qquad b_k = \frac{1}{\sum_x P(k|x)} \sum_x P(k|x)|x - \mu_k|$$

- In a mixture of Gaussians and Laplacians, Gaussians use the Gaussian update rules, Laplacians use the Laplacian rule

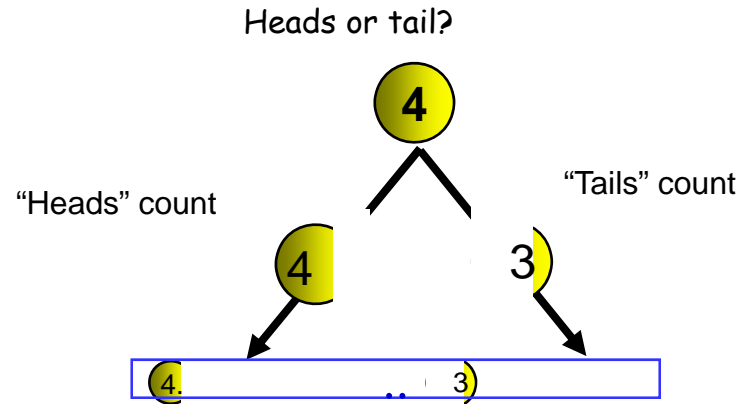
Expectation Maximization

- The EM algorithm is used whenever proper statistical analysis of a phenomenon requires the knowledge of a hidden or missing variable (or a set of hidden/missing variables)
 - The hidden variable is often called a “latent” variable
- Some examples:
 - Estimating mixtures of distributions
 - Only data are observed. The individual distributions and mixing proportions must both be learnt.
 - Estimating the distribution of data, when some attributes are missing
 - Estimating the dynamics of a system, based only on observations that may be a complex function of system state

Solve this problem:

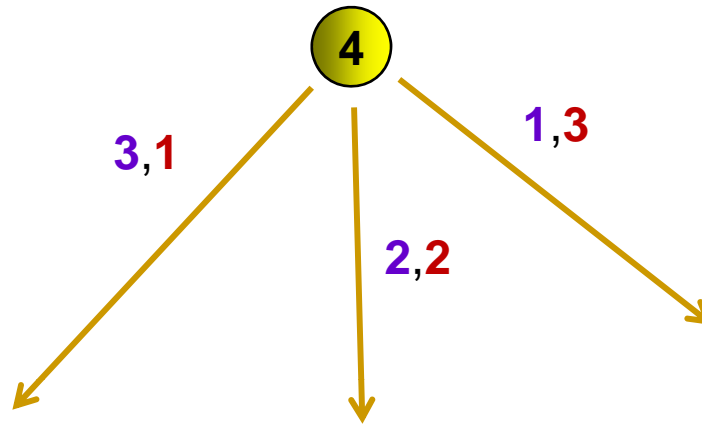
- Caller rolls a dice and flips a coin
 - He calls out the number rolled if the coin shows head
 - Otherwise he calls the number+1
 - Determine $p(\text{heads})$ and $p(\text{number})$ for the dice from a collection of outputs
- Caller rolls two dice
 - He calls out the sum
 - Determine $P(\text{dice})$ from a collection of outputs

The dice and the coin



- Unknown: Whether it was head or tails

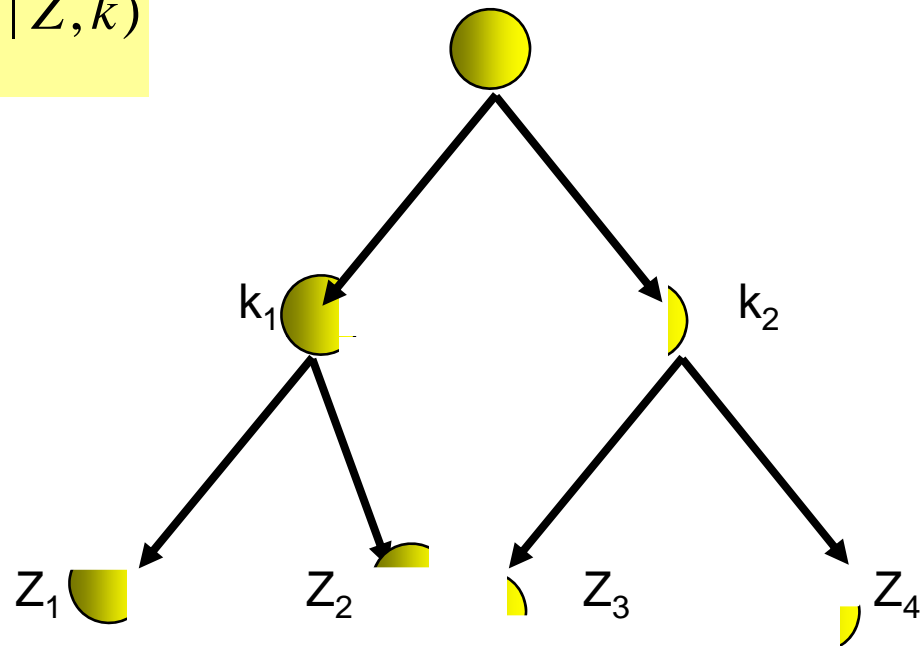
The two dice



- Unknown: How to partition the number
- $\text{Count}_{\text{blue}}(3) += P(3,1 \mid 4)$
- $\text{Count}_{\text{blue}}(2) += P(2,2 \mid 4)$
- $\text{Count}_{\text{blue}}(1) += P(1,3 \mid 4)$

Fragmentation can be hierarchical

$$P(X) = \sum_k P(k) \sum_Z P(Z | k) P(X | Z, k)$$

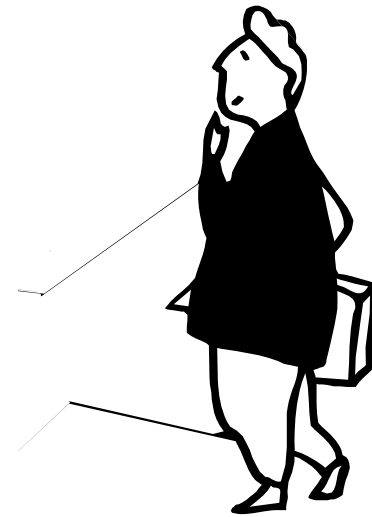


- E.g. mixture of mixtures
- Fragments are further fragmented..
 - Work this out

More later

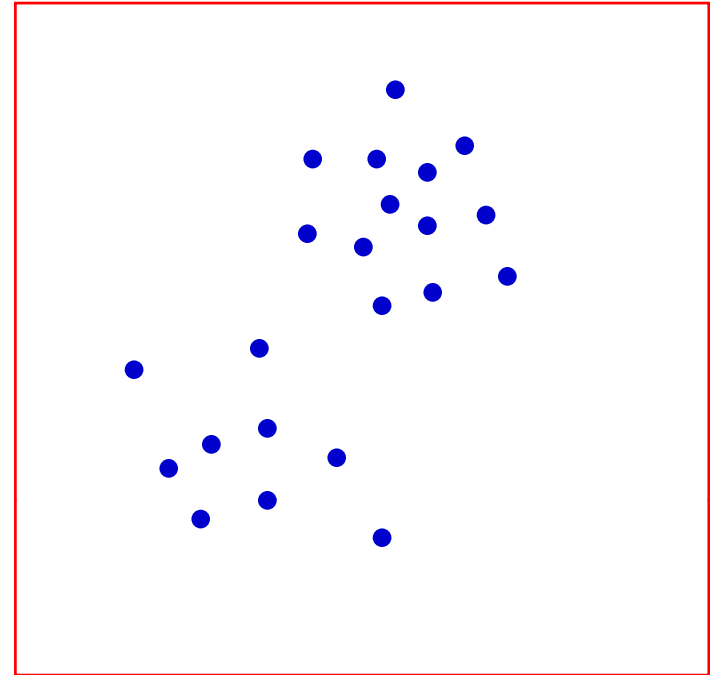
- Will see a couple of other instances of the use of EM
 - E.g. HMM training
 - Homework problems

Clustering



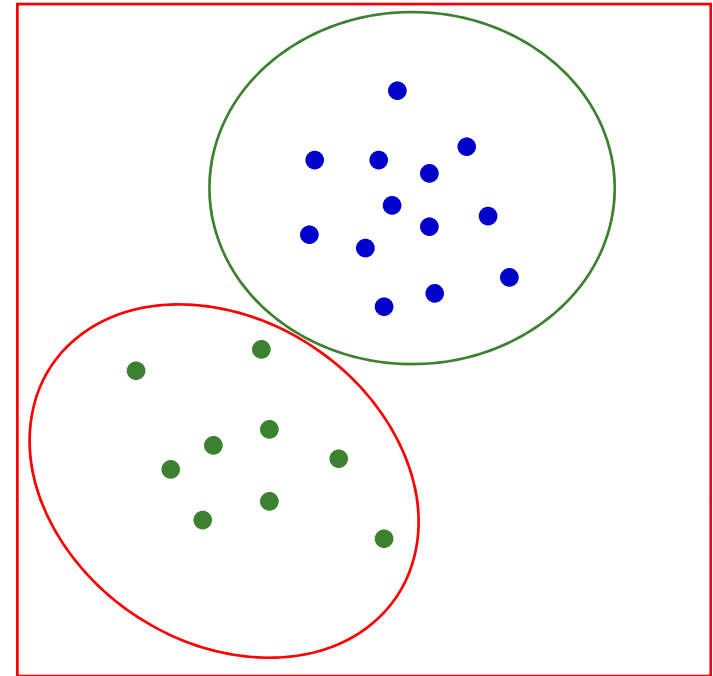
Clustering

- What is clustering
 - Clustering is the determination of naturally occurring grouping of data/instances (with low within-group variability and high between-group variability)



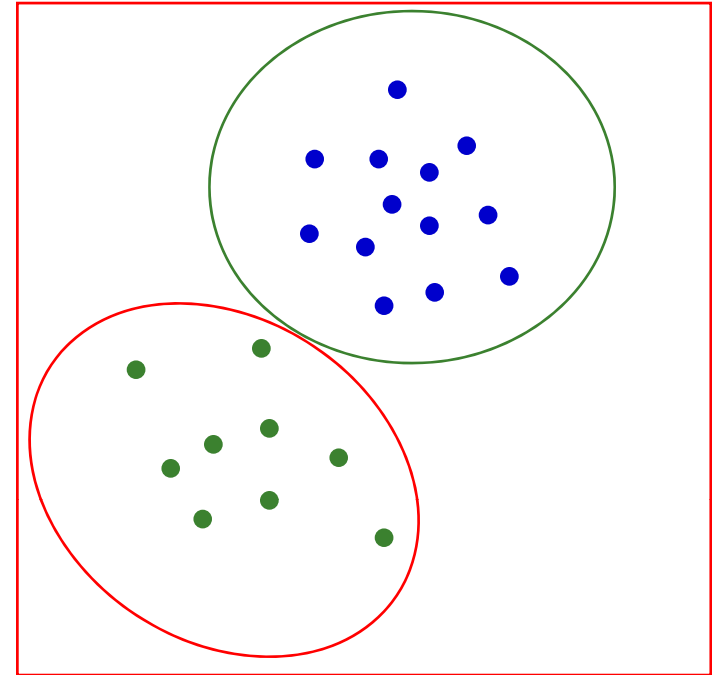
Clustering

- What is clustering
 - Clustering is the determination of naturally occurring grouping of data/instances (with low within-group variability and high between-group variability)



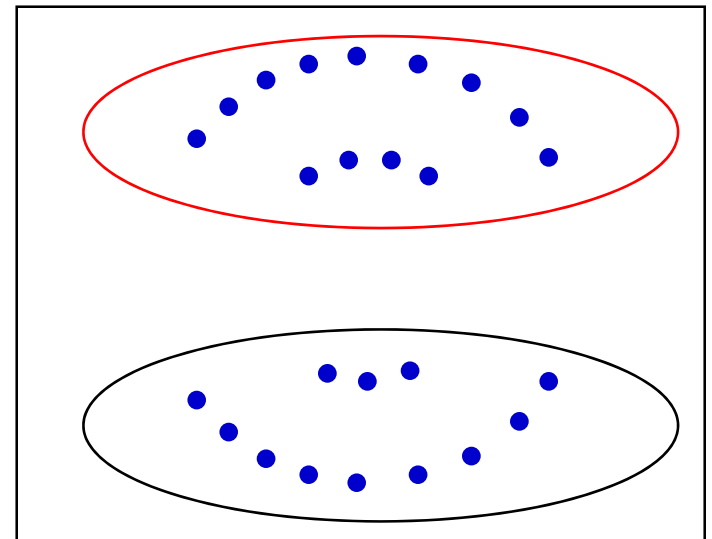
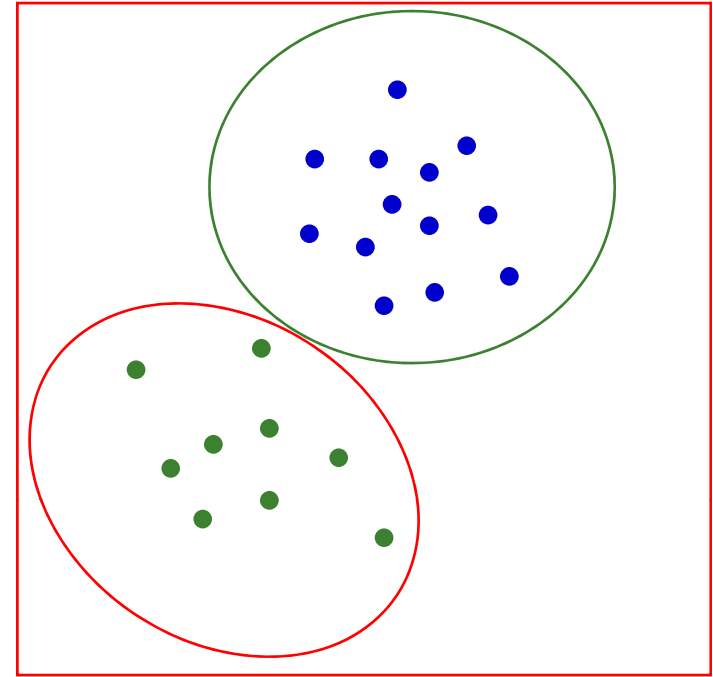
Clustering

- What is clustering
 - Clustering is the determination of naturally occurring grouping of data/instances (with low within-group variability and high between-group variability)
- How is it done
 - Find groupings of data such that the groups optimize a “within-group-variability” objective function of some kind



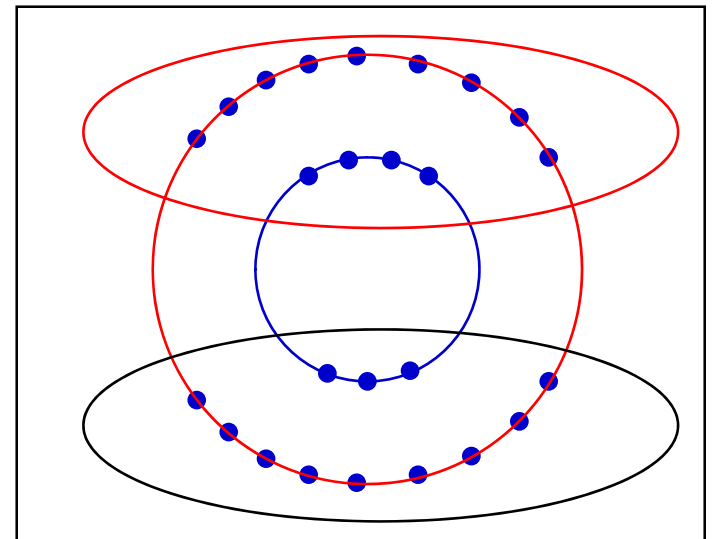
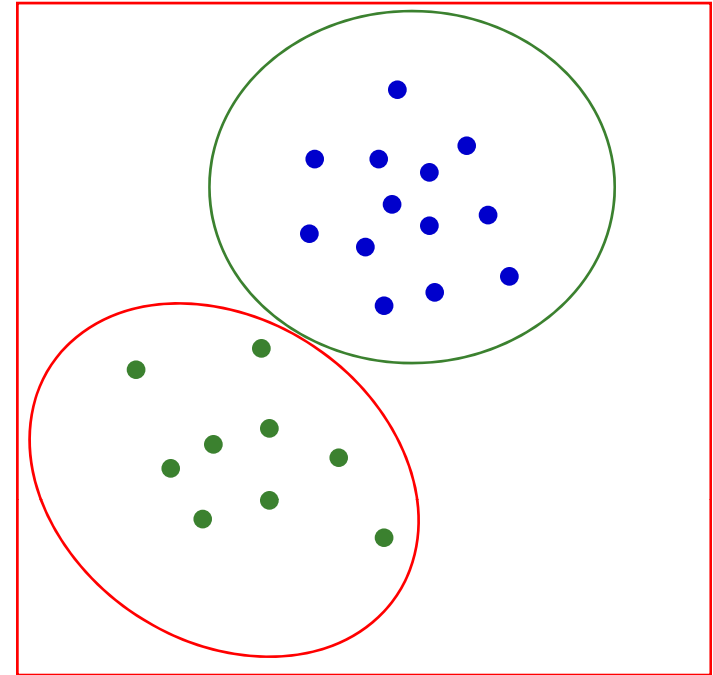
Clustering

- What is clustering
 - Clustering is the determination of naturally occurring grouping of data/instances (with low within-group variability and high between-group variability)
- How is it done
 - Find groupings of data such that the groups optimize a “within-group-variability” objective function of some kind
 - The objective function used affects the nature of the discovered clusters
 - E.g. Euclidean distance and distance from center result in different clusters in this example



Clustering

- What is clustering
 - Clustering is the determination of naturally occurring grouping of data/instances (with low within-group variability and high between-group variability)
- How is it done
 - Find groupings of data such that the groups optimize a “within-group-variability” objective function of some kind
 - The objective function used affects the nature of the discovered clusters
 - E.g. Euclidean distance and distance from center result in different clusters in this example



Why Clustering

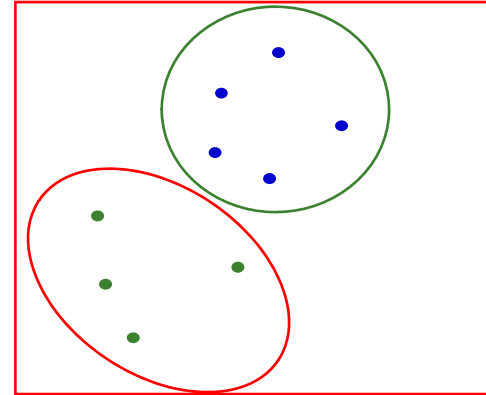
- Automatic grouping into “Classes”
 - Different clusters may show different behavior
- Quantization
 - All data within a cluster are represented by a single point
- Preprocessing step for other algorithms
 - Indexing, categorization, etc.

Clustering criteria

- Compactness criterion
 - Measure that shows how “good” clusters are
 - The objective function
- Distance of a point from a cluster
 - To determine the cluster a data vector belongs to

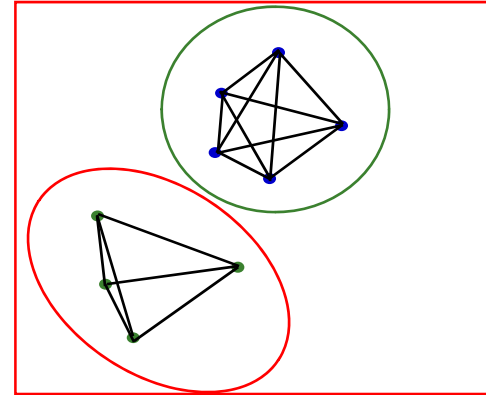
“Compactness” criteria for clustering

- Distance based measures
 - Total distance between each element in the cluster and every other element in the cluster



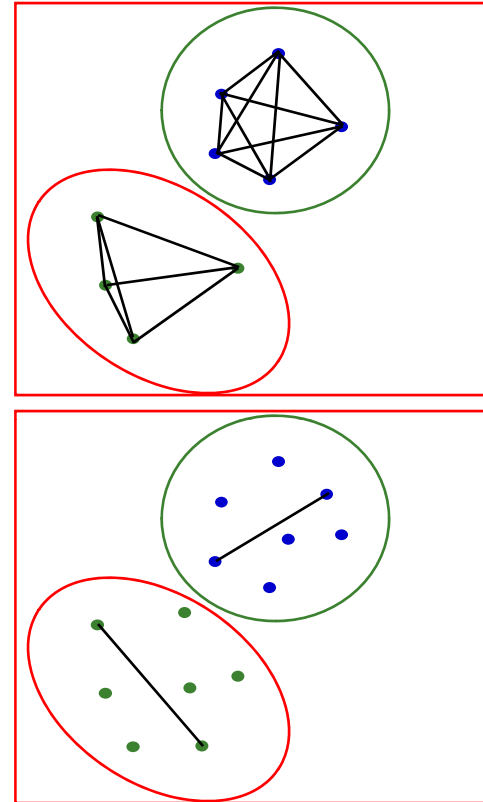
“Compactness” criteria for clustering

- Distance based measures
 - Total distance between each element in the cluster and every other element in the cluster



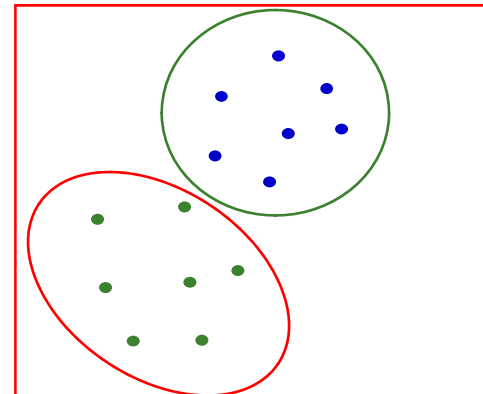
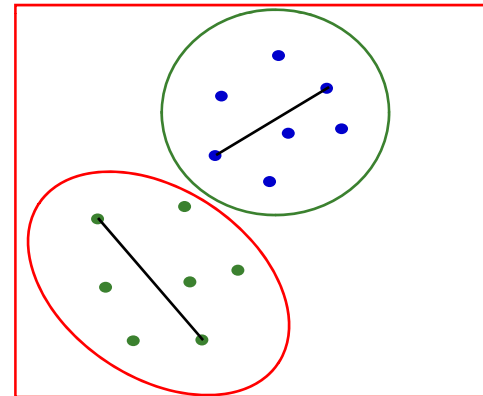
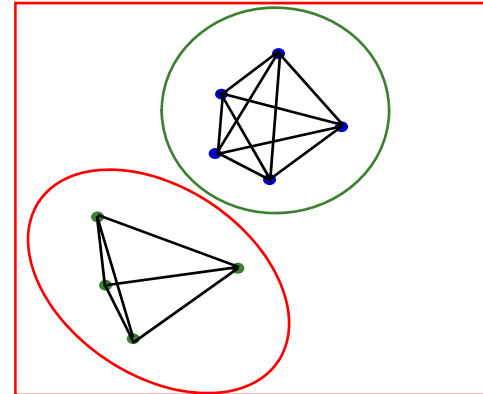
“Compactness” criteria for clustering

- Distance based measures
 - Total distance between each element in the cluster and every other element in the cluster
 - Distance between the two farthest points in the cluster



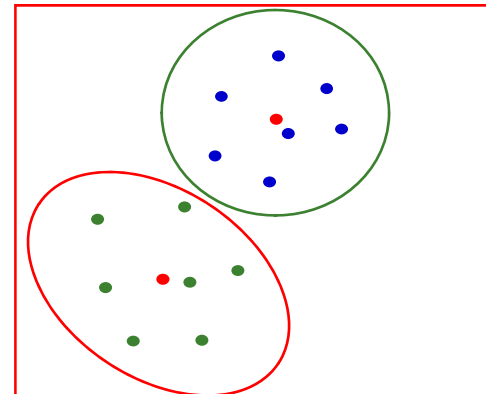
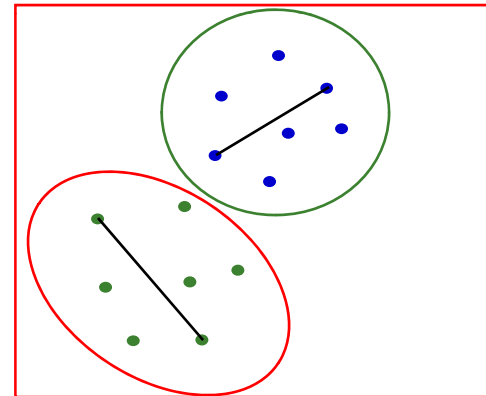
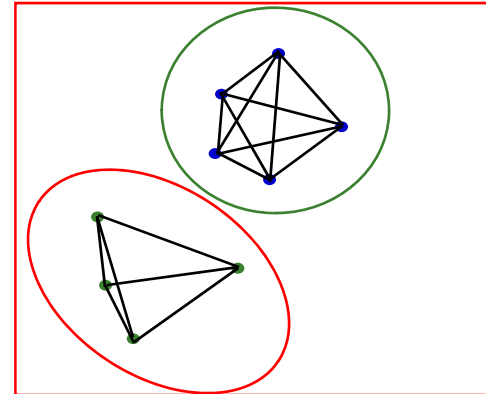
“Compactness” criteria for clustering

- Distance based measures
 - Total distance between each element in the cluster and every other element in the cluster
 - Distance between the two farthest points in the cluster
 - Total distance of every element in the cluster from the centroid of the cluster



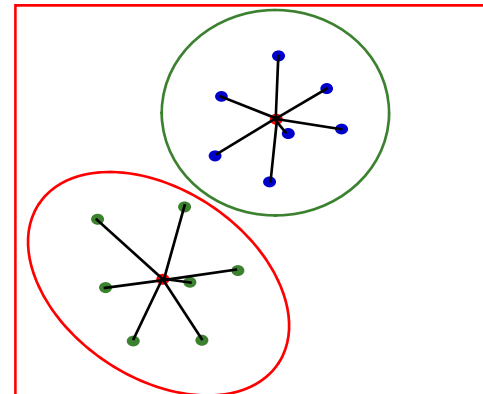
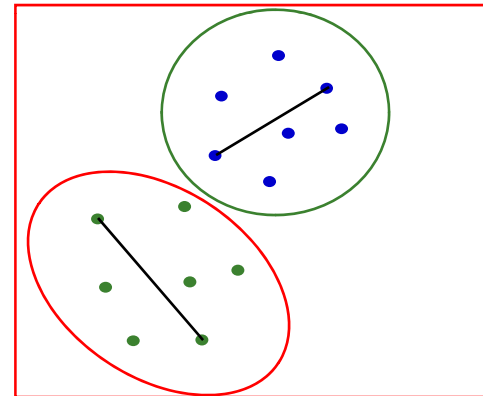
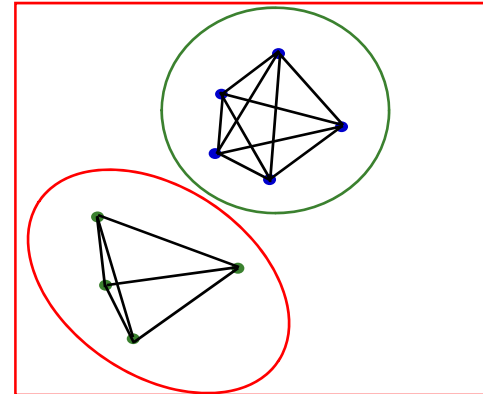
“Compactness” criteria for clustering

- Distance based measures
 - Total distance between each element in the cluster and every other element in the cluster
 - Distance between the two farthest points in the cluster
 - Total distance of every element in the cluster from the centroid of the cluster



“Compactness” criteria for clustering

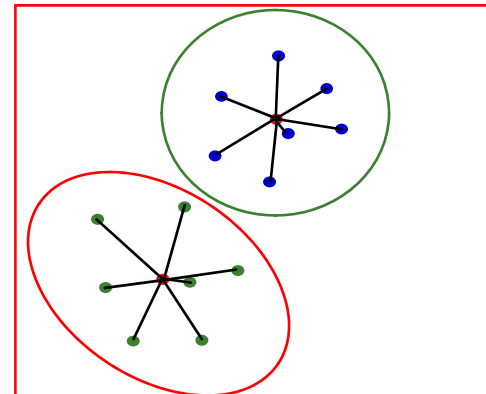
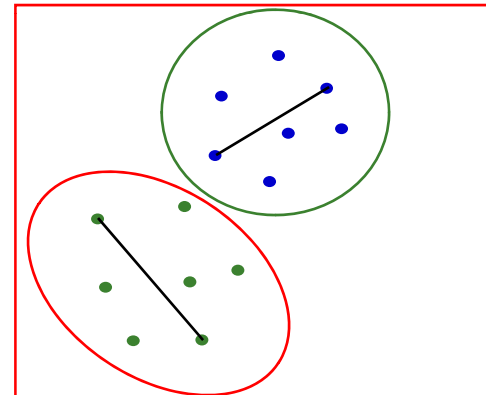
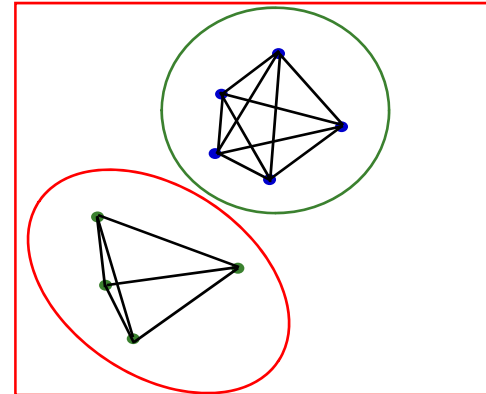
- Distance based measures
 - Total distance between each element in the cluster and every other element in the cluster
 - Distance between the two farthest points in the cluster
 - Total distance of every element in the cluster from the centroid of the cluster



“Compactness” criteria for clustering

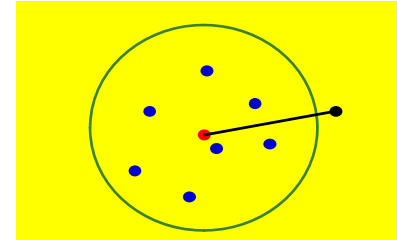
- Distance based measures
 - Total distance between each element in the cluster and every other element in the cluster
 - Distance between the two farthest points in the cluster
 - Total distance of every element in the cluster from the centroid of the cluster
 - Distance measures are often weighted Minkowski metrics

$$dist = \sqrt[n]{w_1|a_1 - b_1|^n + w_2|a_2 - b_2|^n + \dots + w_M|a_M - b_M|^n}$$



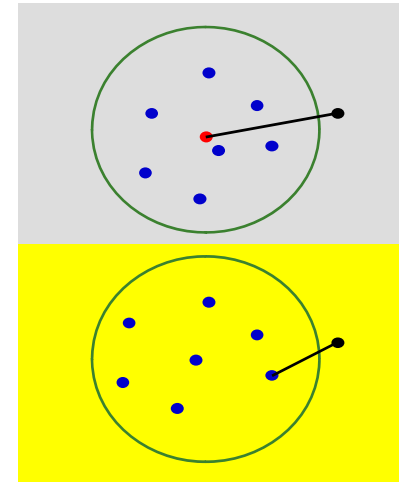
Clustering: Distance from cluster

- How far is a data point from a cluster?
 - Euclidean or Minkowski distance from the centroid of the cluster



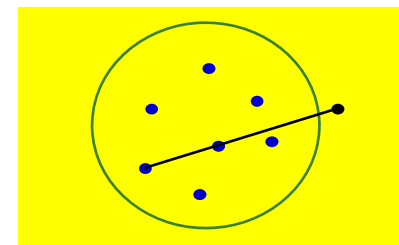
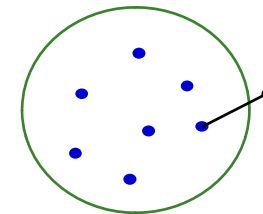
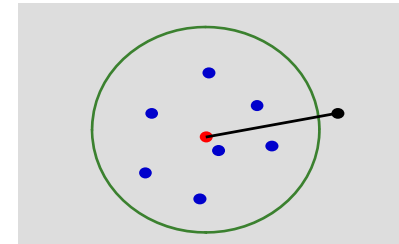
Clustering: Distance from cluster

- How far is a data point from a cluster?
 - Euclidean or Minkowski distance from the centroid of the cluster
 - Distance from the closest point in the cluster



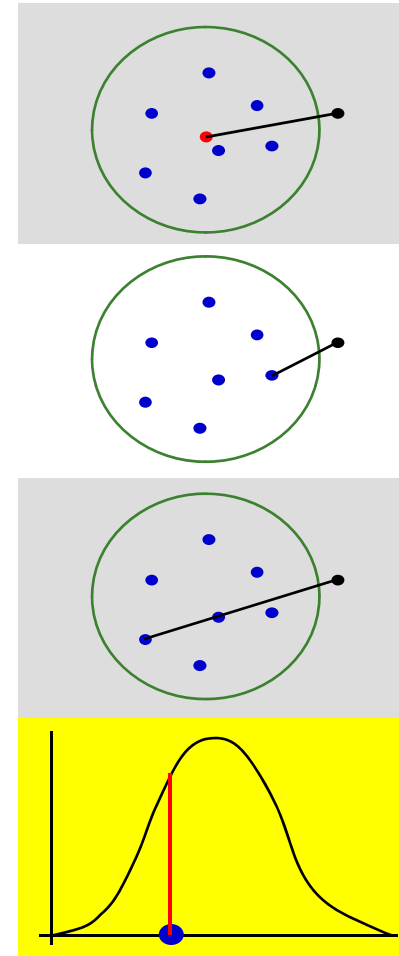
Clustering: Distance from cluster

- How far is a data point from a cluster?
 - Euclidean or Minkowski distance from the centroid of the cluster
 - Distance from the closest point in the cluster
 - Distance from the farthest point in the cluster



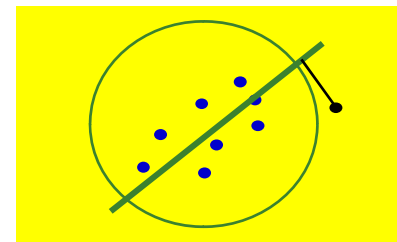
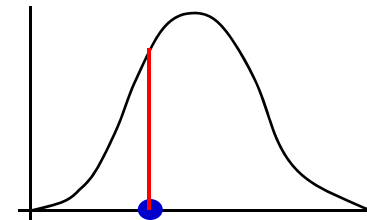
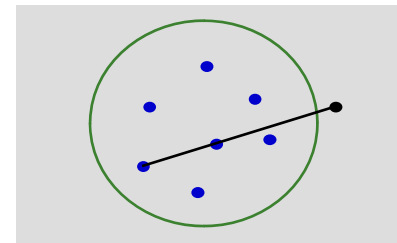
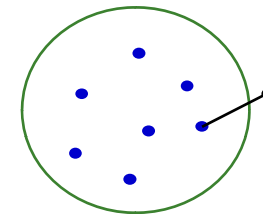
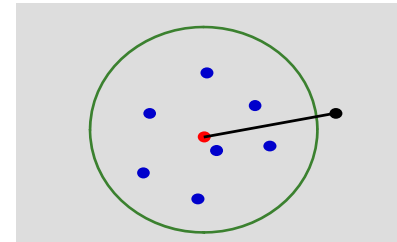
Clustering: Distance from cluster

- How far is a data point from a cluster?
 - Euclidean or Minkowski distance from the centroid of the cluster
 - Distance from the closest point in the cluster
 - Distance from the farthest point in the cluster
 - Probability of data measured on cluster distribution



Clustering: Distance from cluster

- How far is a data point from a cluster?
 - ❑ Euclidean or Minkowski distance from the centroid of the cluster
 - ❑ Distance from the closest point in the cluster
 - ❑ Distance from the farthest point in the cluster
 - ❑ Probability of data measured on cluster distribution
 - ❑ Fit of data to cluster-based regression



Optimal clustering: Exhaustive enumeration

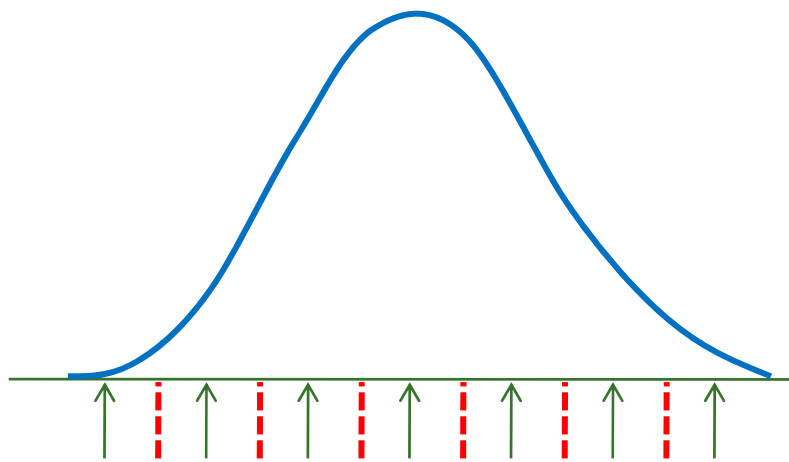
- All possible combinations of data must be evaluated
 - If there are M data points, and we desire N clusters, the number of ways of separating M instances into N clusters is

$$\frac{1}{M!} \sum_{i=0}^N (-1)^i \binom{N}{i} (N-i)^M$$

- Exhaustive enumeration based clustering requires that the objective function (the “Goodness measure”) be evaluated for every one of these, and the best one chosen
- This is the only correct way of optimal clustering
 - Unfortunately, it is also computationally unrealistic

Not-quite non sequitur: Quantization

Probability of analog value

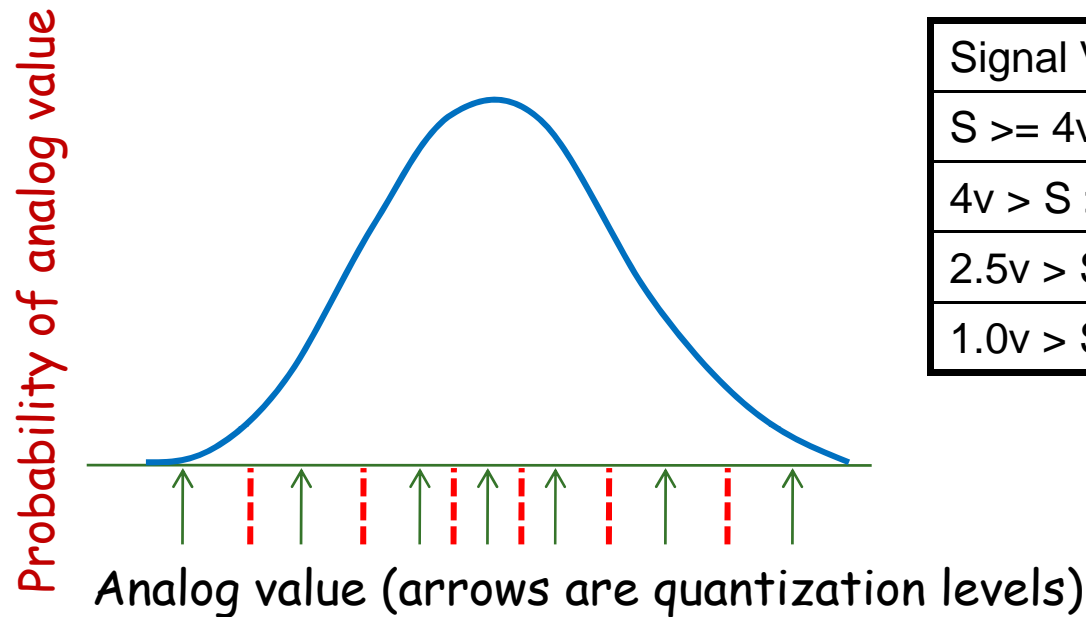


Analog value (arrows are quantization levels)

Signal Value	Bits	Mapped to
$S \geq 3.75v$	11	3 * const
$3.75v > S \geq 2.5v$	10	2 * const
$2.5v > S \geq 1.25v$	01	1 * const
$1.25v > S \geq 0v$	0	0

- Linear quantization (uniform quantization):
 - Each digital value represents an equally wide range of analog values
 - Regardless of distribution of data
 - Digital-to-analog conversion represented by a “uniform” table

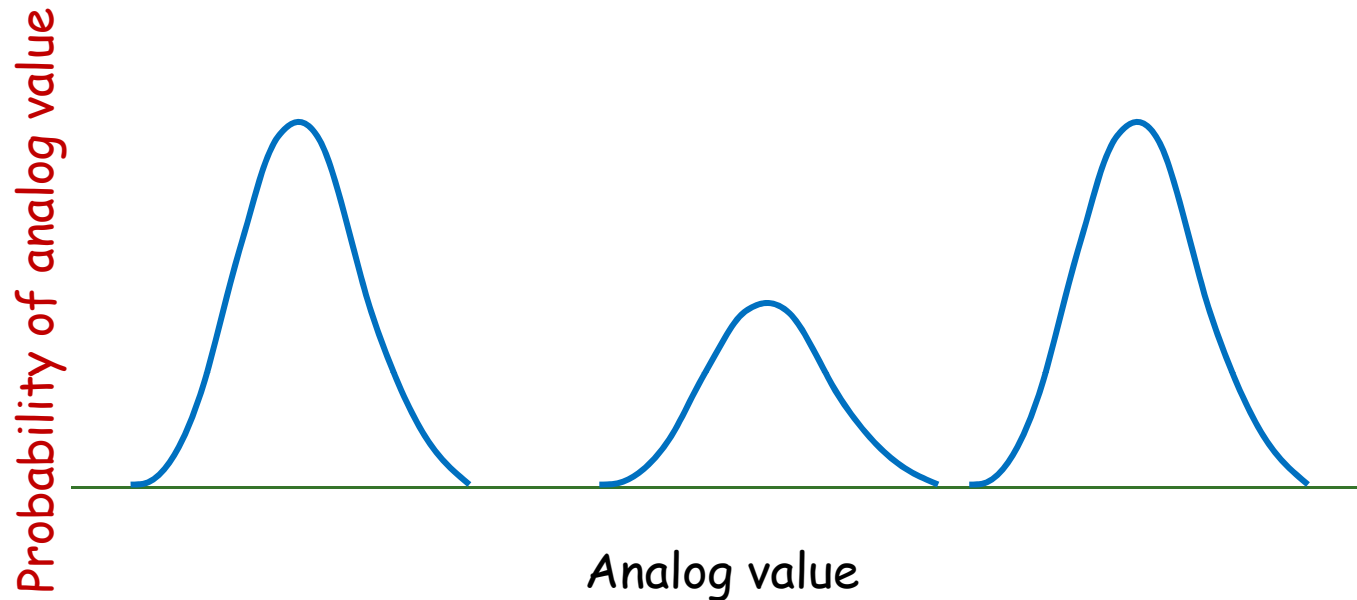
Not-quite non sequitur: Quantization



Signal Value	Bits	Mapped to
$S \geq 4v$	11	4.5
$4v > S \geq 2.5v$	10	3.25
$2.5v > S \geq 1v$	01	1.25
$1.0v > S \geq 0v$	0	0.5

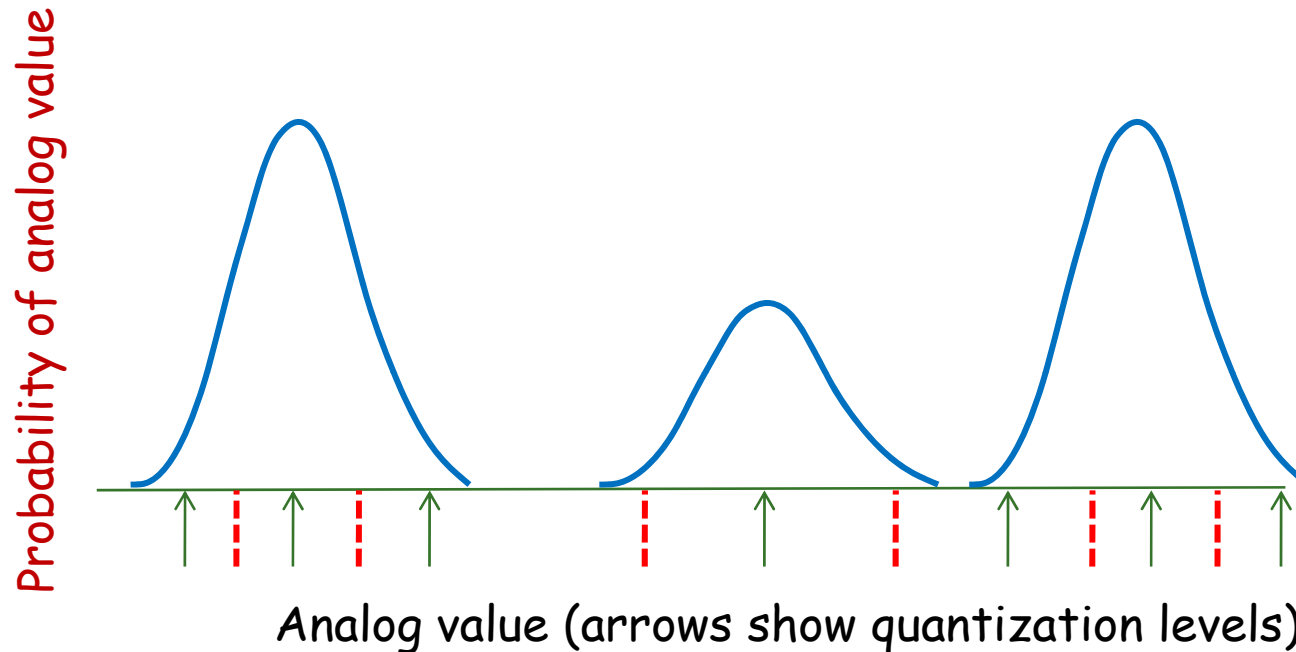
- Non-Linear quantization:
 - Each digital value represents a different range of analog values
 - Finer resolution in high-density areas
 - Mu-law / A-law assumes a gaussian-like distribution of data
 - Digital-to-analog conversion represented by a “non-uniform” table

Non-uniform quantization



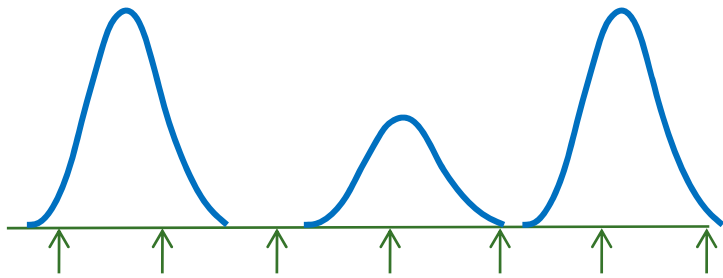
- If data distribution is not Gaussianish?
 - Mu-law / A-law are not optimal
 - How to compute the optimal ranges for quantization
 - Or the optimal table

The Lloyd Quantizer



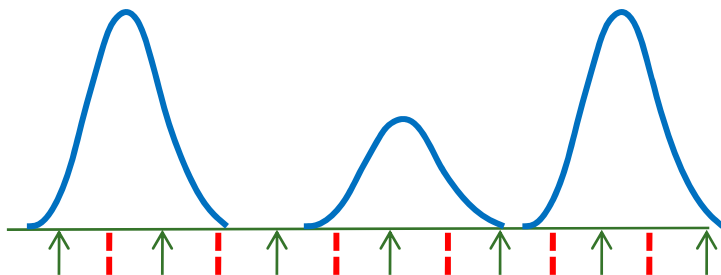
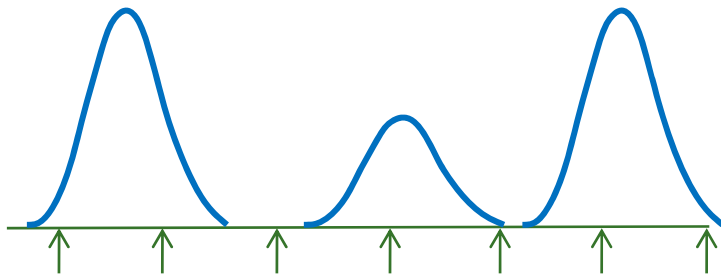
- Lloyd quantizer: An iterative algorithm for computing optimal quantization tables for non-uniformly distributed data
- **Learned from “training” data**

Lloyd Quantizer



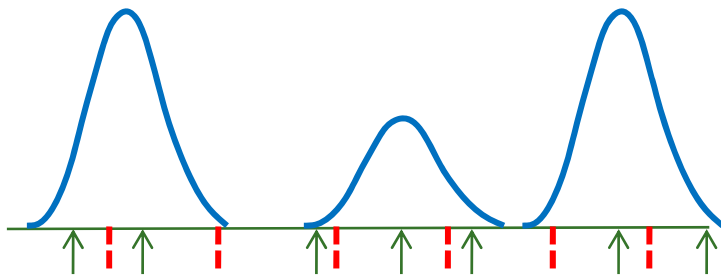
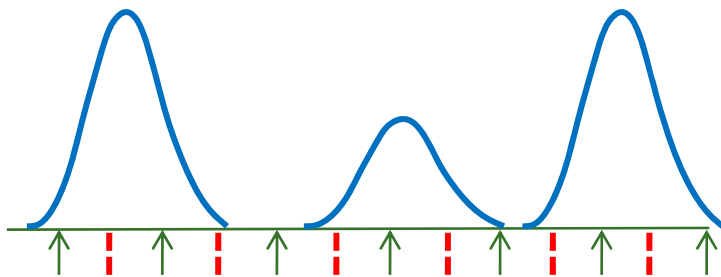
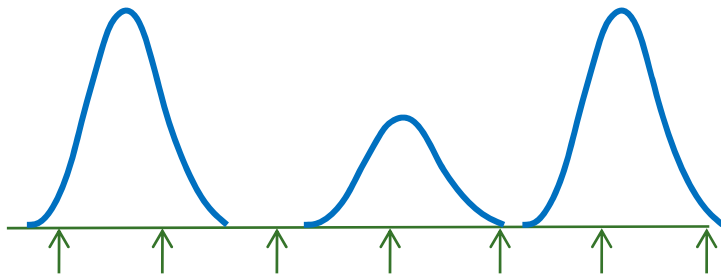
- Randomly initialize quantization points
 - Right column entries of quantization table

Lloyd Quantizer



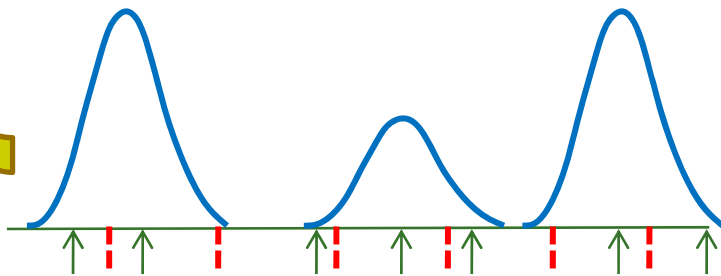
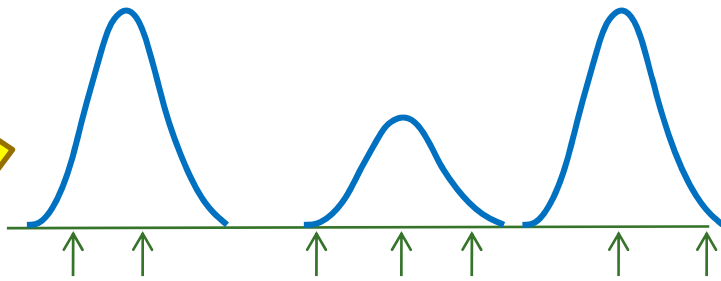
- Randomly initialize quantization points
 - Right column entries of quantization table
- Assign all training points to the nearest quantization point
 - Draw boundaries

Lloyd Quantizer



- Randomly initialize quantization points
 - Right column entries of quantization table
- Assign all training points to the nearest quantization point
 - Draw boundaries
- Reestimate quantization points

Lloyd Quantizer



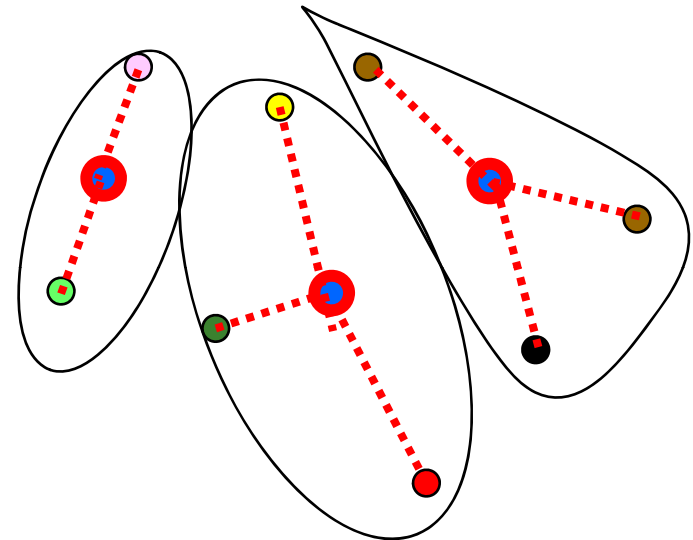
- Randomly initialize quantization points
 - Right column entries of quantization table
- Assign all training points to the nearest quantization point
 - Draw boundaries
- Reestimate quantization points
- Iterate until convergence

Generalized Lloyd Algorithm: K-means clustering

- K means is an iterative algorithm for clustering **vector** data
 - McQueen, J. 1967. "Some methods for classification and analysis of multivariate observations." Proceedings of the Fifth Berkeley Symposium on Mathematical Statistics and Probability, 281-297
- General procedure:
 - Initially group data into the required number of clusters somehow (initialization)
 - Assign each data point to the closest cluster
 - Once all data points are assigned to clusters, redefine clusters
 - Iterate

K-means

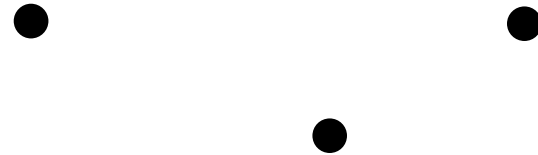
- Problem: Given a set of data vectors, find natural clusters
- Clustering criterion is **scatter**: distance from the centroid
 - Every cluster has a centroid
 - The centroid represents the cluster
- **Definition:** The **centroid** is the weighted mean of the cluster
 - Weight = 1 for basic scheme



$$m_{cluster} = \frac{1}{\sum_{i \in cluster} w_i} \sum_{i \in cluster} w_i x_i$$

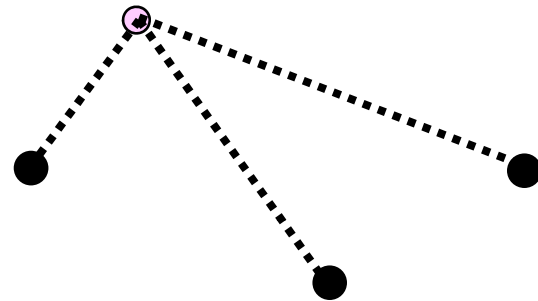
K-means

1. Initialize a set of centroids randomly



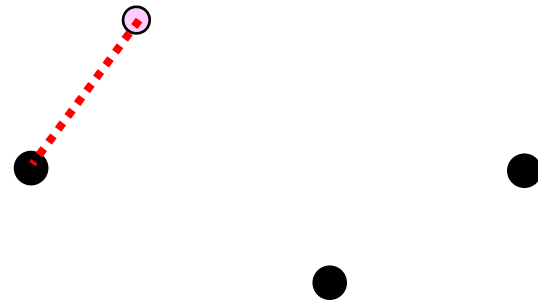
K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$



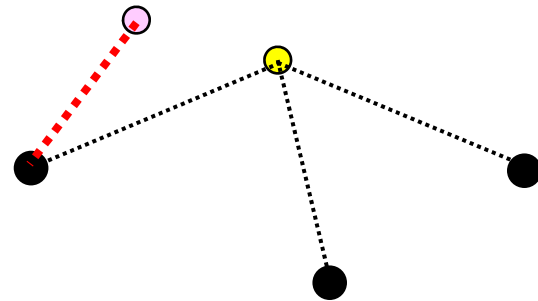
K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum



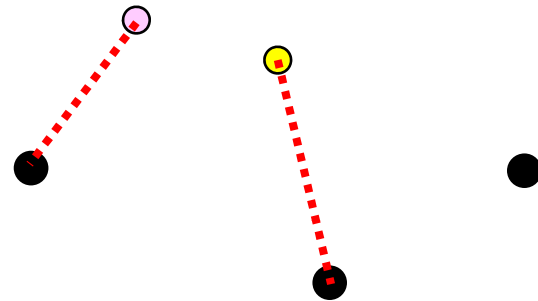
K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum



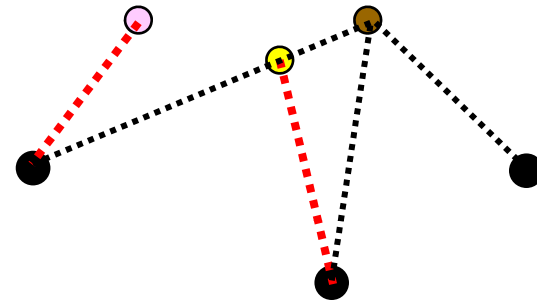
K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum



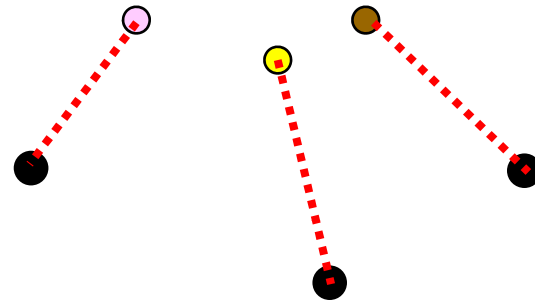
K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum



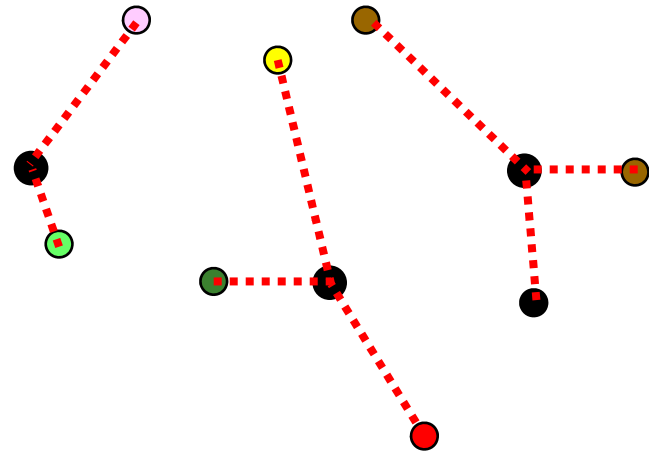
K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum



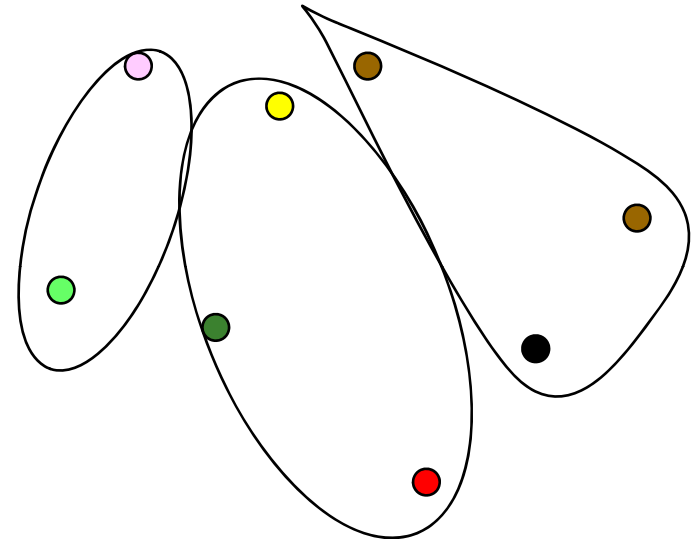
K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum



K-means

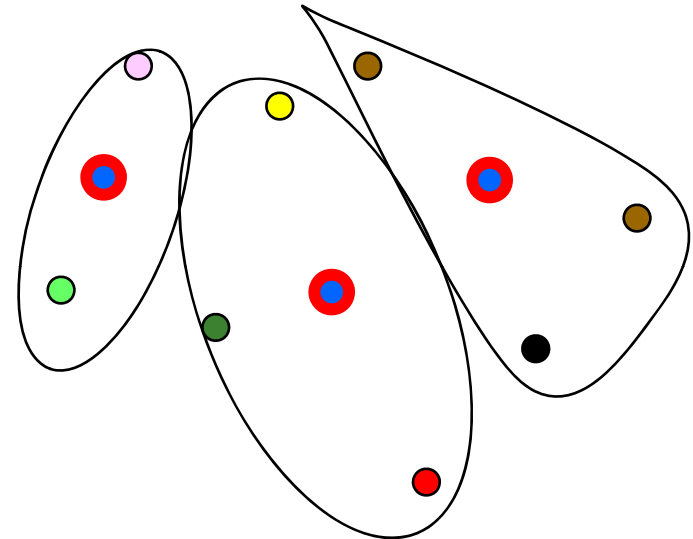
1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum



K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum
4. When all data points are clustered, recompute centroids

$$m_{cluster} = \frac{1}{\sum_{i \in cluster} w_i} \sum_{i \in cluster} w_i x_i$$

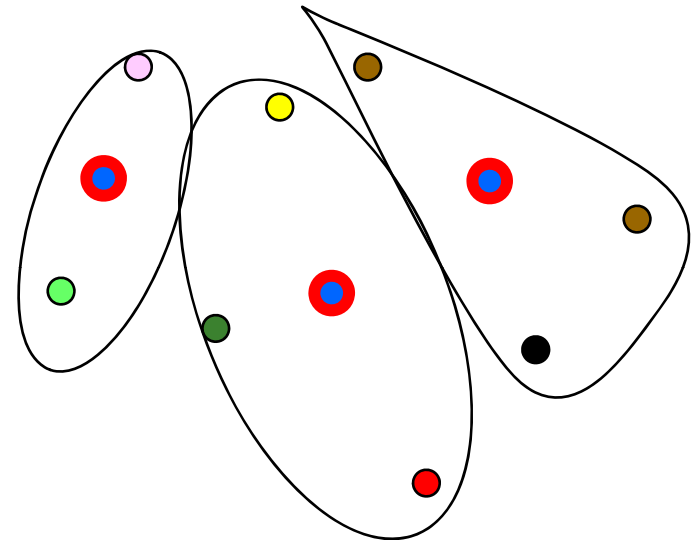


K-means

1. Initialize a set of centroids randomly
2. For each data point x , find the distance from the centroid for each cluster
 - $d_{cluster} = \mathbf{distance}(x, m_{cluster})$
3. Put data point in the cluster of the closest centroid
 - Cluster for which $d_{cluster}$ is minimum
4. When all data points are clustered, recompute centroids

$$m_{cluster} = \frac{1}{\sum_{i \in cluster} w_i} \sum_{i \in cluster} w_i x_i$$

5. If not converged, go back to 2



K-Means comments

- The distance metric determines the clusters
 - In the original formulation, the distance is L2 distance
 - Euclidean norm, $w_i = 1$

$$\text{distance}_{cluster}(x, m_{cluster}) = \|x - m_{cluster}\|_2$$

$$m_{cluster} = \frac{1}{N_{cluster}} \sum_{i \in cluster} x_i$$

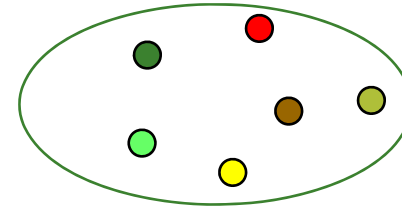
- If we replace every x by $m_{cluster}(x)$, we get *Vector Quantization*
- K-means is an instance of *generalized EM*
- Not guaranteed to converge for all distance metrics

Initialization

- Random initialization
- Top-down clustering
 - Initially partition the data into two (or a small number of) clusters using K means
 - Partition each of the resulting clusters into two (or a small number of) clusters, also using K means
 - Terminate when the desired number of clusters is obtained

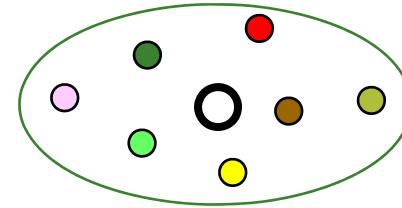
K-Means for Top-Down clustering

1. Start with one cluster



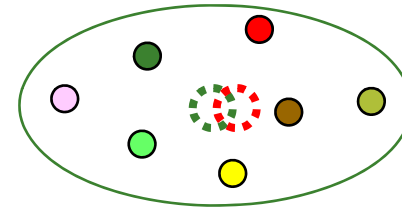
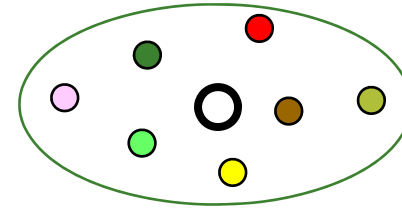
K-Means for Top-Down clustering

1. Start with one cluster
2. Split each cluster into two:
 - Perturb centroid of cluster slightly (by $< 5\%$) to generate two centroids



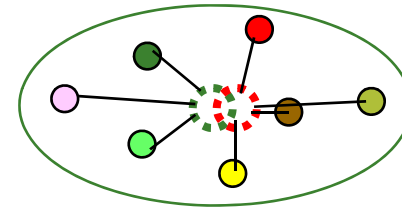
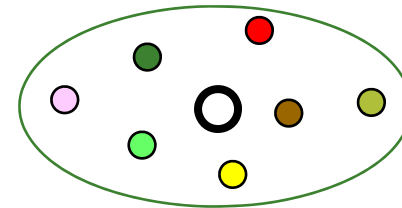
K-Means for Top-Down clustering

1. Start with one cluster
2. Split each cluster into two:
 - Perturb centroid of cluster slightly (by $< 5\%$) to generate two centroids



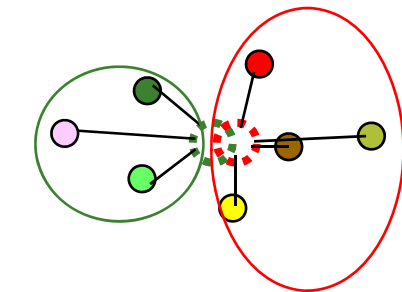
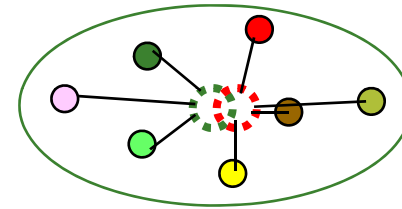
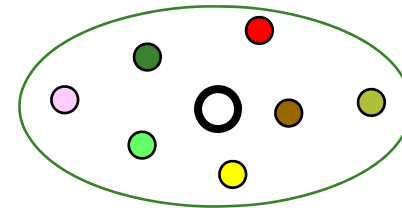
K-Means for Top-Down clustering

1. Start with one cluster
2. Split each cluster into two:
 - Perturb centroid of cluster slightly (by $< 5\%$) to generate two centroids
3. Initialize K means with new set of centroids



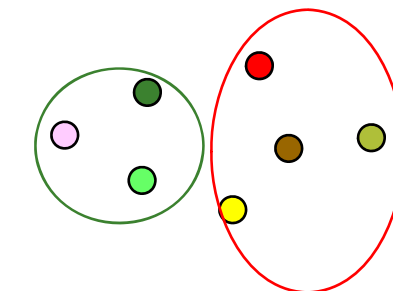
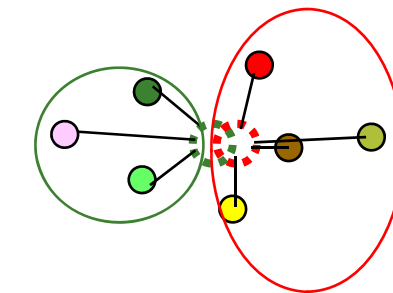
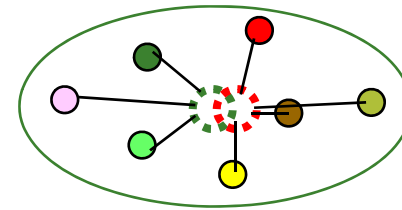
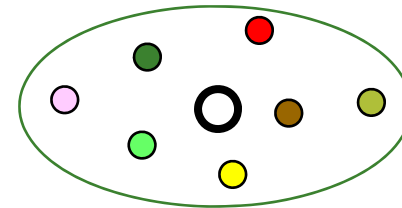
K-Means for Top-Down clustering

1. Start with one cluster
2. Split each cluster into two:
 - Perturb centroid of cluster slightly (by $< 5\%$) to generate two centroids
3. Initialize K means with new set of centroids
4. Iterate Kmeans until convergence



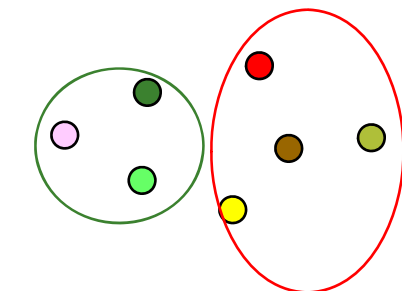
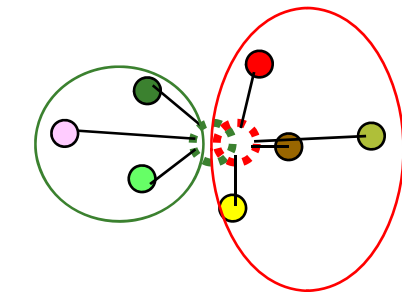
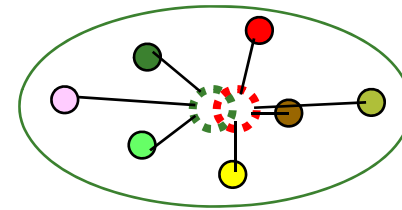
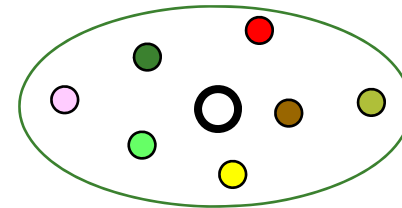
K-Means for Top-Down clustering

1. Start with one cluster
2. Split each cluster into two:
 - Perturb centroid of cluster slightly (by $< 5\%$) to generate two centroids
3. Initialize K means with new set of centroids
4. Iterate Kmeans until convergence

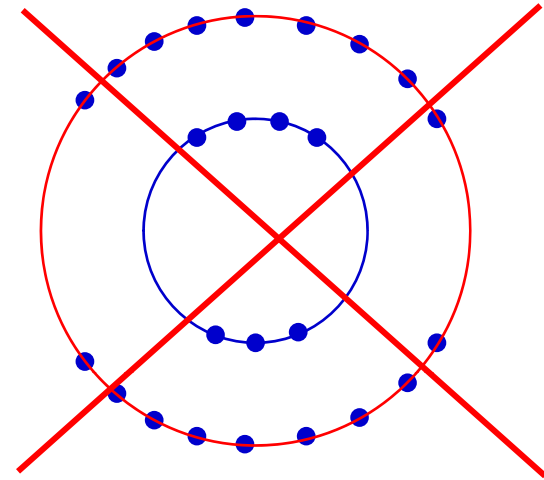
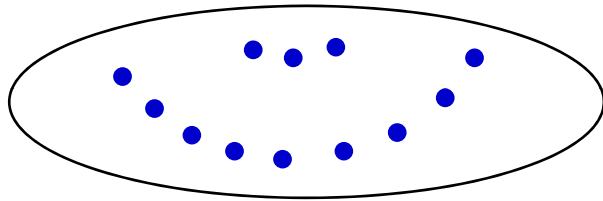
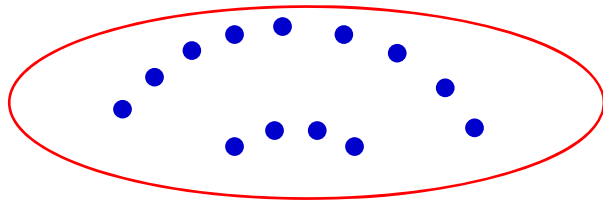


K-Means for Top-Down clustering

1. Start with one cluster
2. Split each cluster into two:
 - Perturb centroid of cluster slightly (by $< 5\%$) to generate two centroids
3. Initialize K means with new set of centroids
4. Iterate Kmeans until convergence
5. If the desired number of clusters is not obtained, return to 2

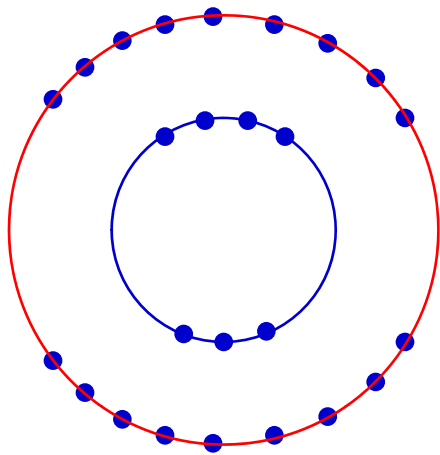


K-means, distances, kernels and spectra

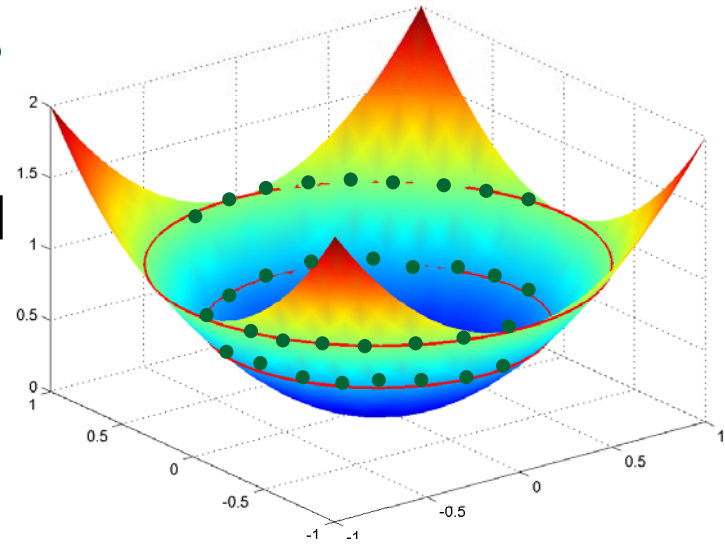


- Basic K-means results in good clusters in Euclidean spaces
 - Alternately stated, will only find clusters that are “good” in terms of Euclidean distances
- Will not find other types of clusters

Non-euclidean clusters



$$f([x,y]) \rightarrow [x,y,z]$$
$$x = x$$
$$y = y$$
$$z = x^2 + y^2$$



- For other forms of clusters we must modify the distance measure
 - E.g. distance from a circle
- May be viewed as a distance in a higher dimensional space
 - I.e *Kernel* distances
 - *Kernel* K-means
- Other related clustering mechanisms:
 - Spectral clustering
 - Non-linear weighting of adjacency
 - Normalized cuts..