

CSE 427
Autumn 2021
MLE, EM

Outline

HW#1 Discussion

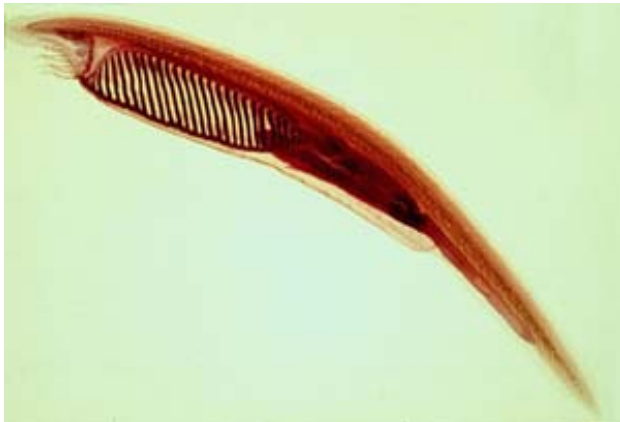
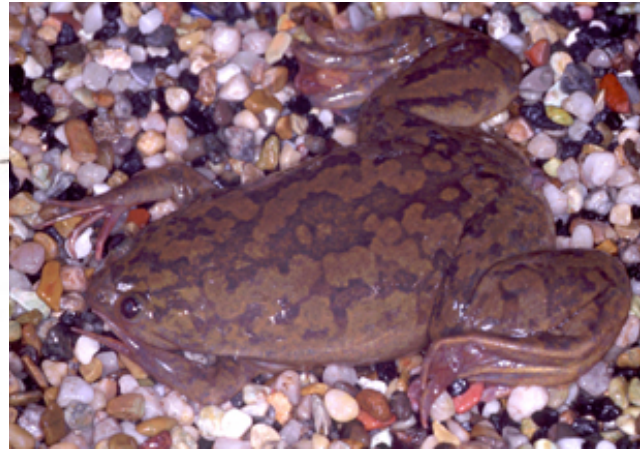
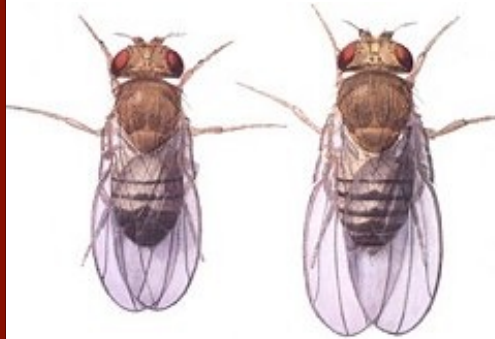
MLE: Maximum Likelihood Estimators

EM: the Expectation Maximization Algorithm

Next: Motif description & discovery

HW # 1 Discussion

	Species	Name	Description	Accession	score to #1
1	Homo sapiens (Human)	MYOD1_HUMAN	Myoblast determination protein 1	P15172	1709
2	Homo sapiens (Human)	TALI_HUMAN	T-cell acute lymphocytic leukemia protein 1 (TAL-1)	P17542	143
3	Mus musculus (Mouse)	MYOD1_MOUSE	Myoblast determination protein 1	P10085	1500
4	Gallus gallus (Chicken)	MYOD1_CHICK	Myoblast determination protein 1 homolog (MYOD1 homolog)	P16075	1020
5	Xenopus laevis (African clawed frog)	MYODA_XENLA	Myoblast determination protein 1 homolog A (Myogenic factor 1)	P13904	978
6	Danio rerio (Zebrafish)	MYOD1_DANRE	Myoblast determination protein 1 homolog (Myogenic factor 1)	Q90477	893
7	Branchiostoma belcheri (Amphioxus)	Q8IU24_BRABE	MyoD-related	Q8IU24	428
8	Drosophila melanogaster (Fruit fly)	MYOD_DROME	Myogenic-determination protein (Protein nautilus) (dMyd)	P22816	368
9	Caenorhabditis elegans	LIN32_CAEEL	Protein lin-32 (Abnormal cell lineage protein 32)	Q10574	118
10	Homo sapiens (Human)	SYFM_HUMAN	Phenylalanyl-tRNA synthetase, mitochondrial	O95363	56



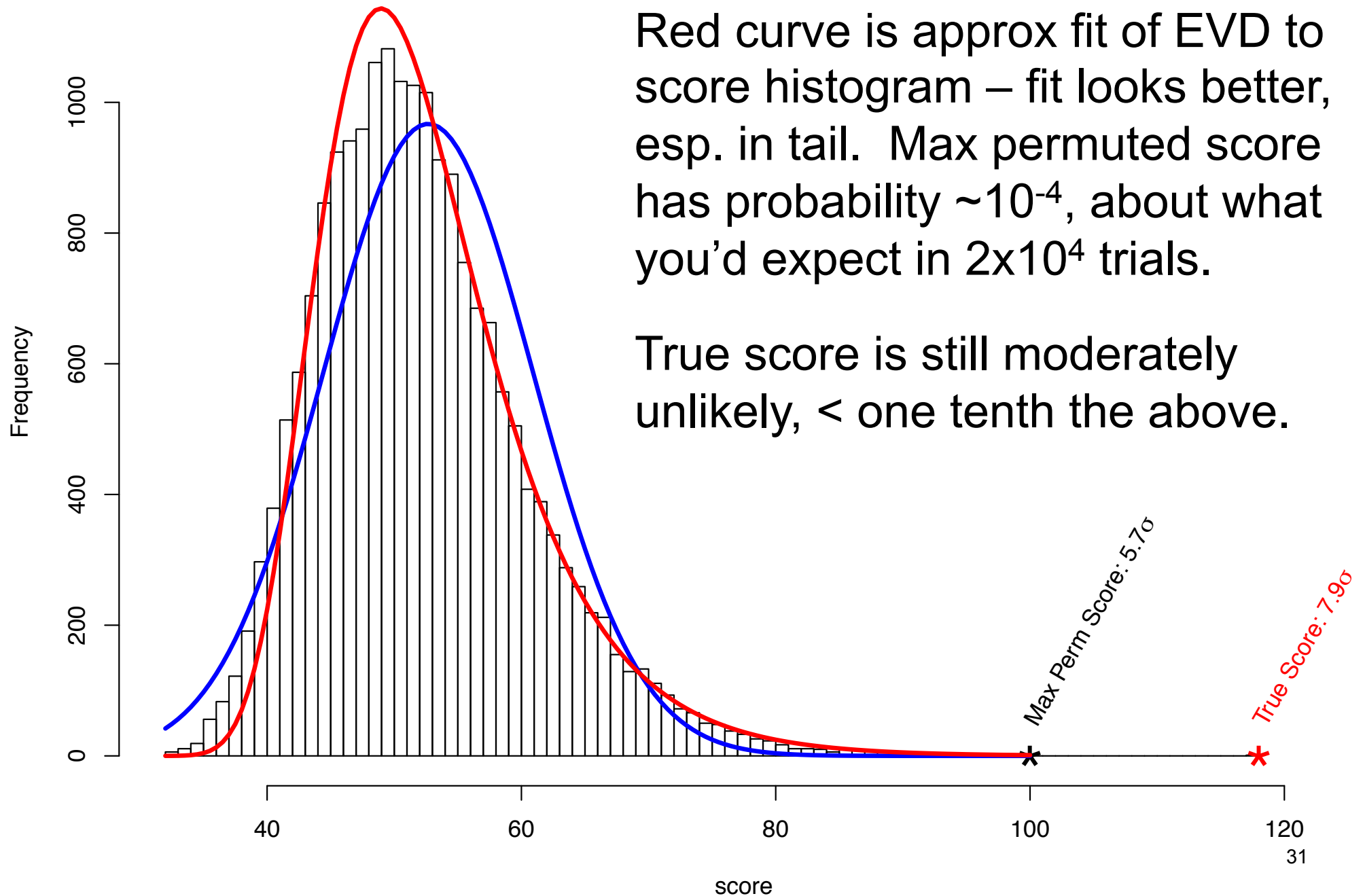
MyoD



jmol_S

<http://www.rcsb.org/pdb/explore/jmol.do?structureId=1MDY&bionumber=1>

Permutation Score Histogram vs Gaussian



Full pairwise score table, reordered

```
## ..... hsMYOD mmMYOD ggMYOD xlMYOD drMYOD bbQ8IU dmMYOD hsTAL1 eLIN32 hsSYFM
```

##	P15172	P10085	P16075	P13904	Q90477	Q8IU24	P22816	P17542	Q10574	O95363
## P15172	1709	1500	1020	978	893	428	368	143	118	56
## P10085	1500	1702	1043	1002	925	440	367	128	118	52
## P16075	1020	1043	1594	1147	1093	448	414	129	120	61
## P13904	978	1002	1147	1541	1104	450	410	128	120	72
## Q90477	893	925	1093	1104	1479	449	410	112	117	62
## Q8IU24	428	440	448	450	449	1215	446	144	125	45
## P22816	368	367	414	410	410	446	1746	123	124	74
## P17542	143	128	129	128	112	144	123	1731	156	66
## Q10574	118	118	120	120	117	125	124	156	746	67
## O95363	56	52	61	72	62	45	74	66	67	2420

species - hs,mm, gg=chick, cl=frog, bb=amphioxus, fly, elegans

Take Home

- Recognizable similarity in protein sequences over great evolutionary distance
- S-W + p-values allow useful quantification
- Similarity correlates better to “same function” than to “same species”

Learning From Data: MLE

Maximum Likelihood Estimators

Parameter Estimation

Given: independent samples x_1, x_2, \dots, x_n from a parametric distribution $f(x|\theta)$

Goal: estimate θ .

Not formally “conditional probability,”
but the notation is convenient...

E.g.: Given sample HHTTTTTHTHTTTHH of (possibly biased) coin flips, estimate

$\theta =$ probability of Heads

$f(x|\theta)$ is the Bernoulli probability mass function with parameter θ

Likelihood

(For *Discrete* Distributions)

$P(x \mid \theta)$: Probability of event x given *model* θ

Viewed as a function of x (fixed θ), it's a *probability*

$$\text{E.g., } \sum_x P(x \mid \theta) = 1$$

Viewed as a function of θ (fixed x), it's called *likelihood*

E.g., $\sum_{\theta} P(x \mid \theta)$ can be anything; *relative* values are the focus.

E.g., if θ = prob of heads in a sequence of coin flips then

$$P(\text{HHTHH} \mid .6) > P(\text{HHTHH} \mid .5),$$

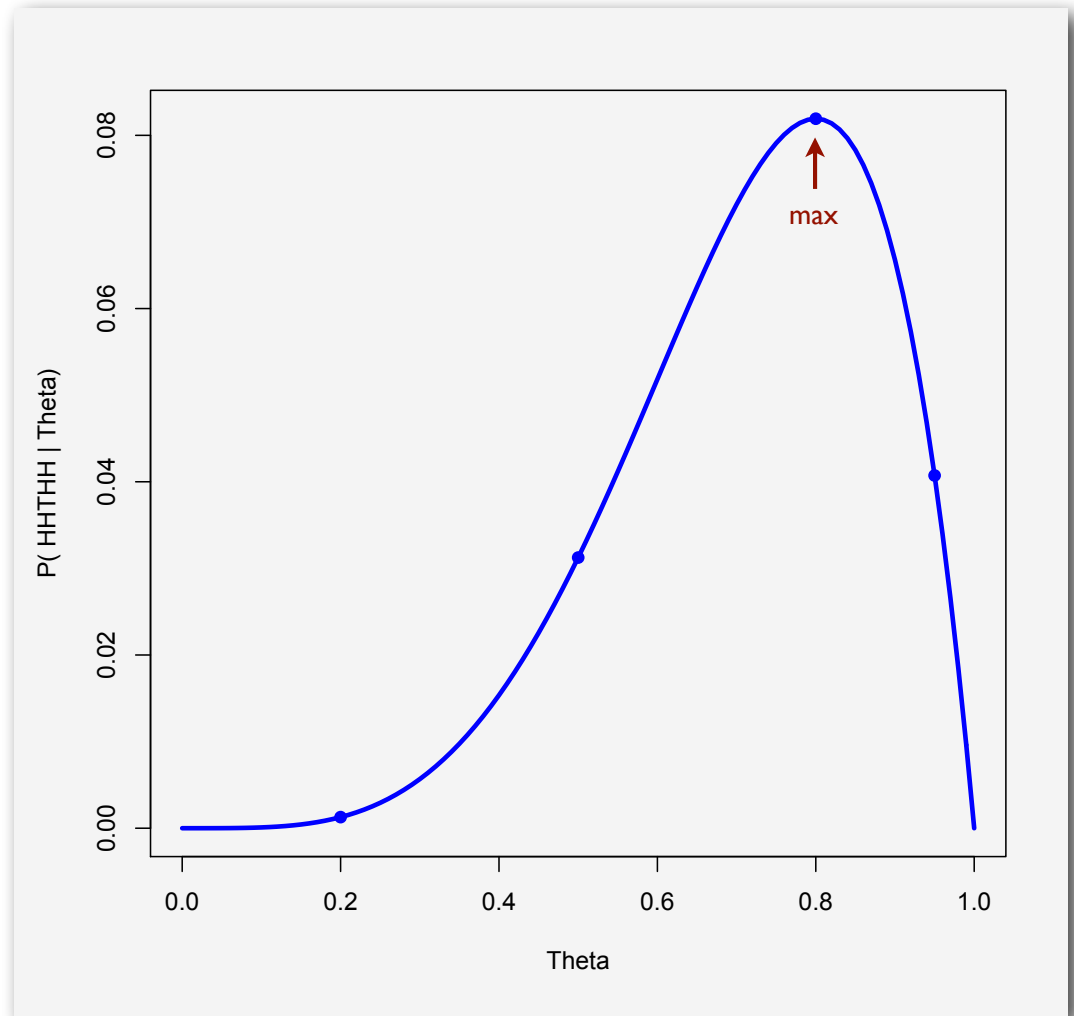
I.e., event HHTHH is *more likely* when $\theta = .6$ than $\theta = .5$

And **what θ make HHTHH *most likely*?**

Likelihood Function

$P(\text{HHTHH} \mid \theta)$:
Probability of HHTHH,
given $P(H) = \theta$:

θ	$\theta^4(1-\theta)$
0.2	0.0013
0.5	0.0313
0.8	0.0819
0.95	0.0407



Maximum Likelihood Parameter Estimation

(For *Discrete* Distributions)

One (of many) approaches to param. est.

Likelihood of (indp) observations x_1, x_2, \dots, x_n

$$L(x_1, x_2, \dots, x_n \mid \theta) = \prod_{i=1}^n f(x_i \mid \theta) \quad (*)$$

As a function of θ , *what θ maximizes the likelihood of the data actually observed?*

Typical approach: $\frac{\partial}{\partial \theta} L(\vec{x} \mid \theta) = 0$ or $\frac{\partial}{\partial \theta} \log L(\vec{x} \mid \theta) = 0$

(*) In general, (discrete) likelihood is the *joint* pmf; product form follows from independence

Example 1

n independent coin flips, x_1, x_2, \dots, x_n ; n_0 tails, n_1 heads,
 $n_0 + n_1 = n$; θ = probability of heads

$$L(x_1, x_2, \dots, x_n \mid \theta) = (1 - \theta)^{n_0} \theta^{n_1}$$

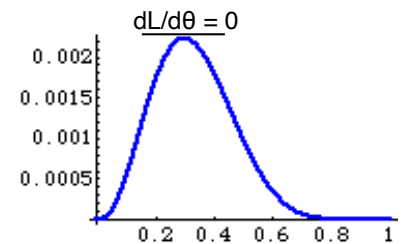
$$\log L(x_1, x_2, \dots, x_n \mid \theta) = n_0 \log(1 - \theta) + n_1 \log \theta$$

$$\frac{\partial}{\partial \theta} \log L(x_1, x_2, \dots, x_n \mid \theta) = \frac{-n_0}{1 - \theta} + \frac{n_1}{\theta}$$

Setting to zero and solving:

$$\hat{\theta} = \frac{n_1}{n}$$

Observed fraction of
successes in *sample* is
MLE of success
probability in *population*



(Also verify it's max, not min, & not better on boundary)

Likelihood

(For *Continuous* Distributions)

$Pr(\text{any specific } x_i) = 0$, so “likelihood = probability” won’t work. Defn: “likelihood” of x_1, \dots, x_n is their *joint density*; = (by indep) product of their *marginal densities*. (As usual, swap *density* for *pmf*.) Why sensible:

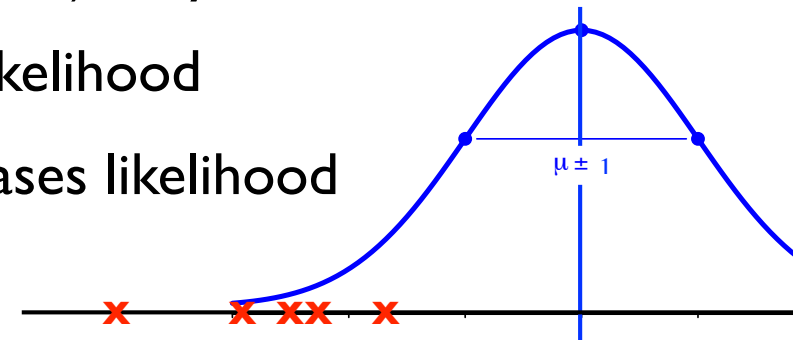
a) density captures all that matters: *relative* likelihood

b) desirable property: better model fit increases likelihood

and

c) if density at x is $f(x)$, for any small $\delta > 0$, the probability of a sample within $\pm \delta/2$ of x is $\approx \delta f(x)$, so density really *is* capturing probability, and δ is *constant* wrt θ , so it just drops out of $d/d\theta \log L(\dots) = 0$.

Otherwise, MLE is just like discrete case: get likelihood, $\frac{\partial}{\partial \theta} \log L(\vec{x} | \theta) = 0$



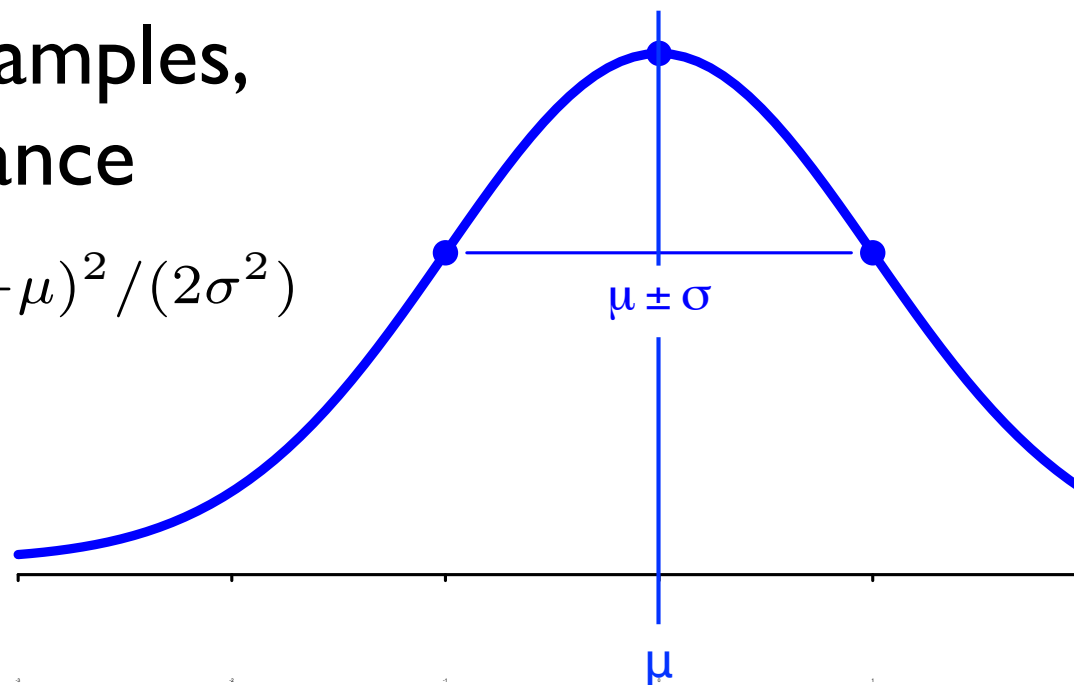
Parameter Estimation

Given: indep samples x_1, x_2, \dots, x_n from a parametric distribution $f(x|\theta)$, **estimate:** θ .

E.g.: Given n normal samples, estimate mean & variance

$$f(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-(x-\mu)^2/(2\sigma^2)}$$

$$\theta = (\mu, \sigma^2)$$



Ex: I got data; a little birdie tells me
it's normal, and promises $\sigma^2 = 1$

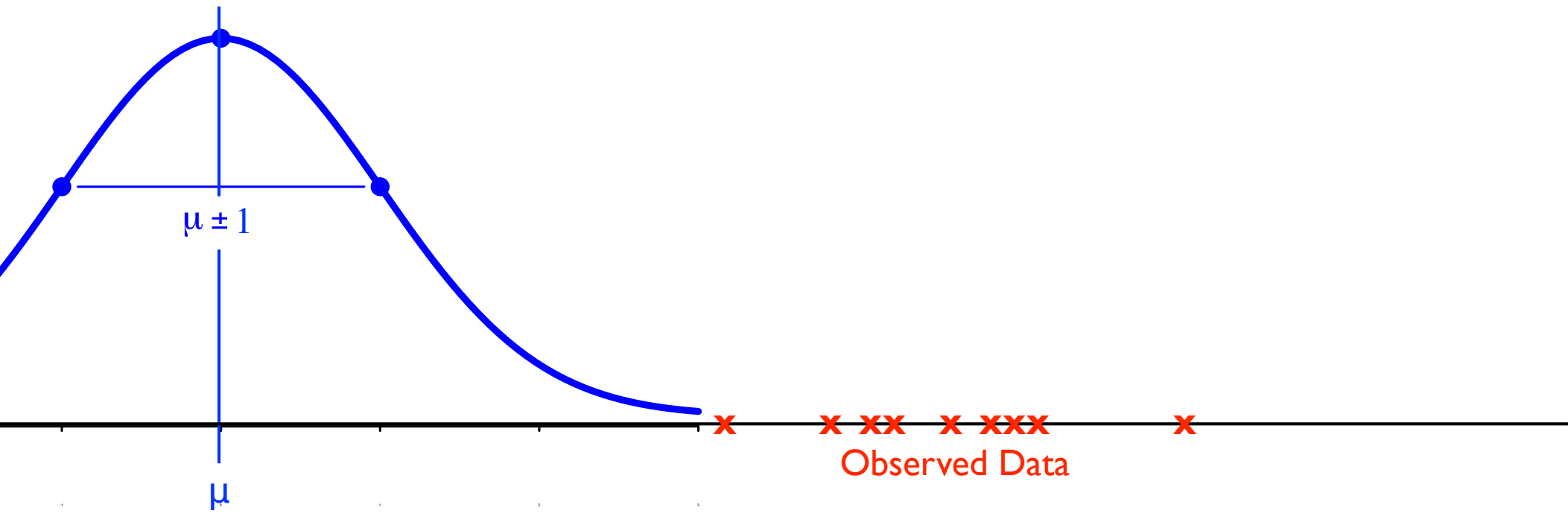


Observed Data

$x \rightarrow$

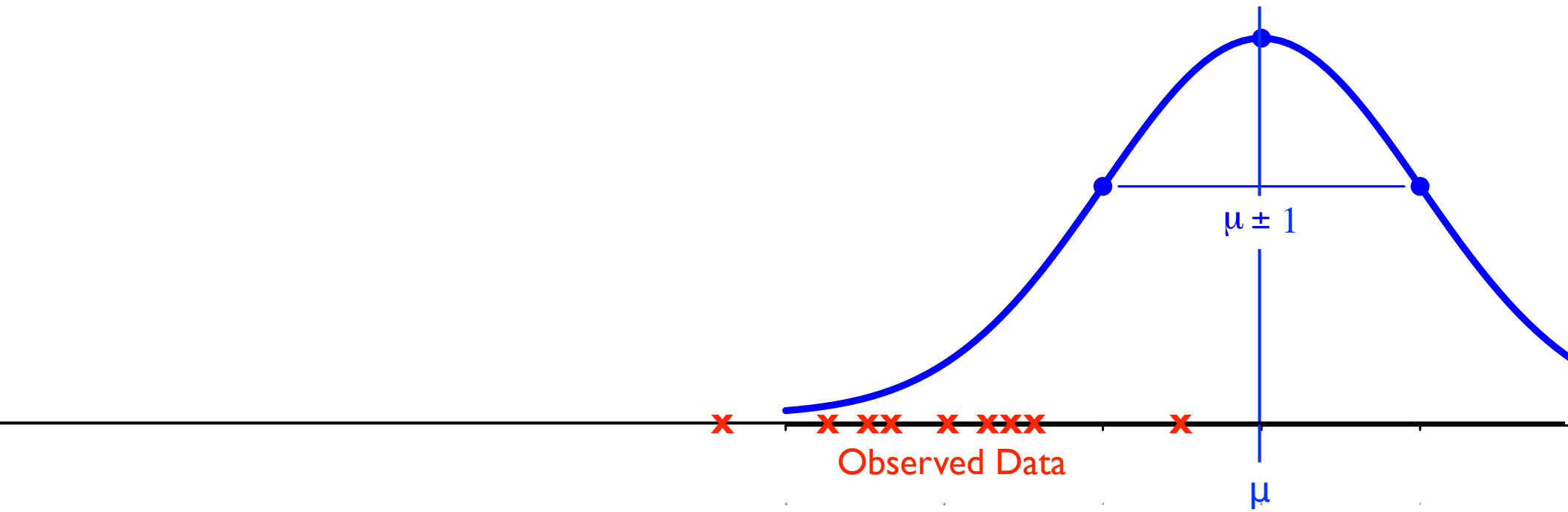
Which is more likely: (a) this?

μ unknown, $\sigma^2 = 1$



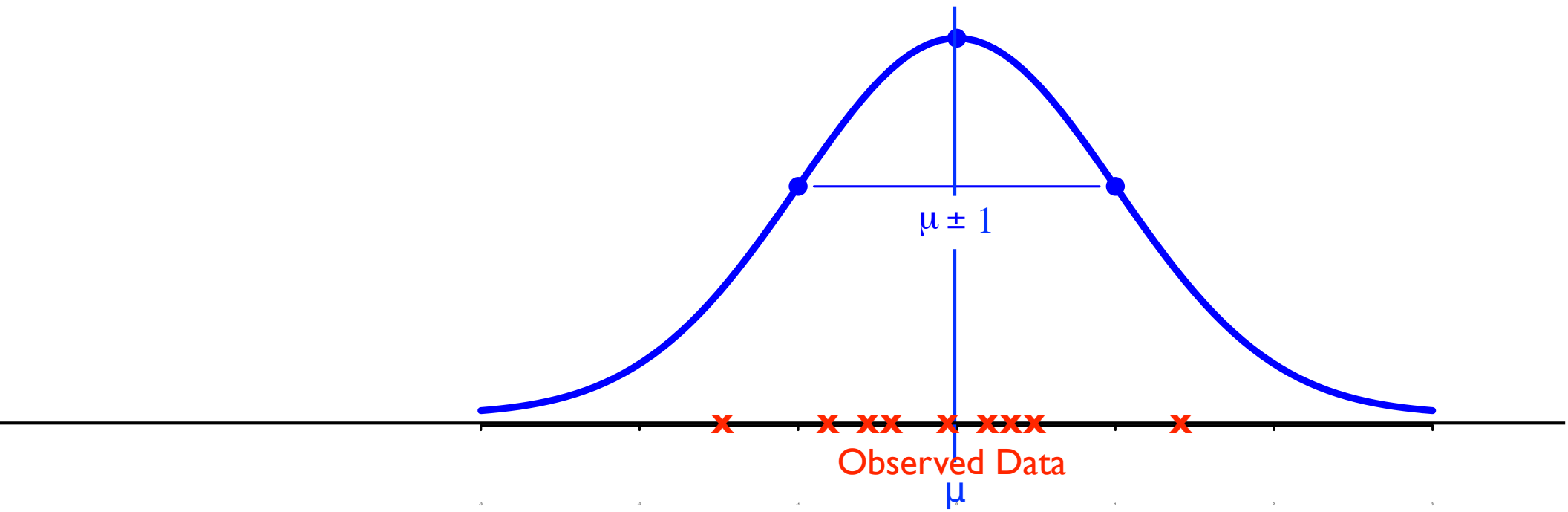
Which is more likely: (b) or this?

μ unknown, $\sigma^2 = 1$



Which is more likely: (c) or *this*?

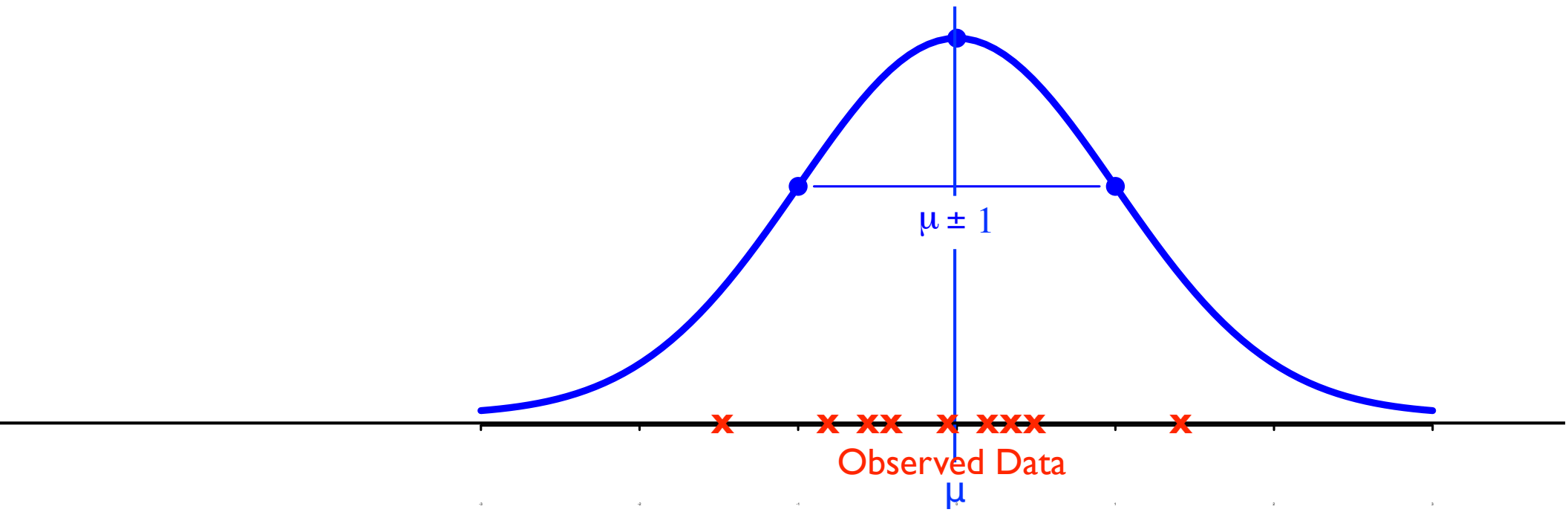
μ unknown, $\sigma^2 = 1$



Which is more likely: (c) or this?

μ unknown, $\sigma^2 = 1$

Looks good by eye, but how do I optimize my estimate of μ ?



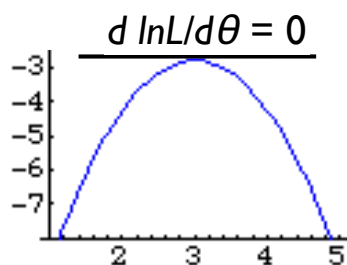
Ex: $x_i \sim N(\mu, \sigma^2), \sigma^2 = 1, \mu$ unknown

$$L(x_1, x_2, \dots, x_n | \theta) = \prod_{i=1}^n \frac{1}{\sqrt{2\pi}} e^{-(x_i - \theta)^2 / 2} \quad \leftarrow \text{product of densities}$$

$$\ln L(x_1, x_2, \dots, x_n | \theta) = \sum_{i=1}^n -\frac{1}{2} \ln(2\pi) - \frac{(x_i - \theta)^2}{2}$$

$$\frac{d}{d\theta} \ln L(x_1, x_2, \dots, x_n | \theta) = \sum_{i=1}^n (x_i - \theta)$$

And verify it's max,
not min & not better
on boundary



$$= \left(\sum_{i=1}^n x_i \right) - n\theta = 0$$

$$\hat{\theta} = \left(\sum_{i=1}^n x_i \right) / n = \bar{x}$$

Sample mean is MLE of
population mean

Ex: I got data; a little birdie tells me it's normal (but does *not* tell me μ , σ^2)

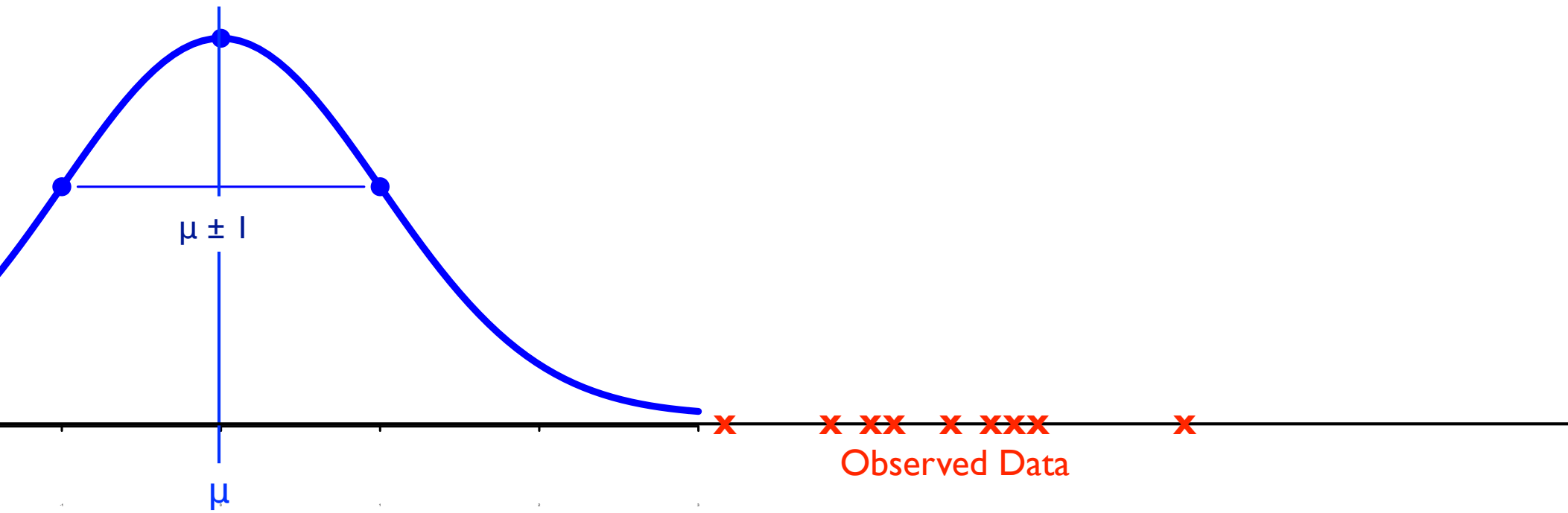


Observed Data

$x \rightarrow$

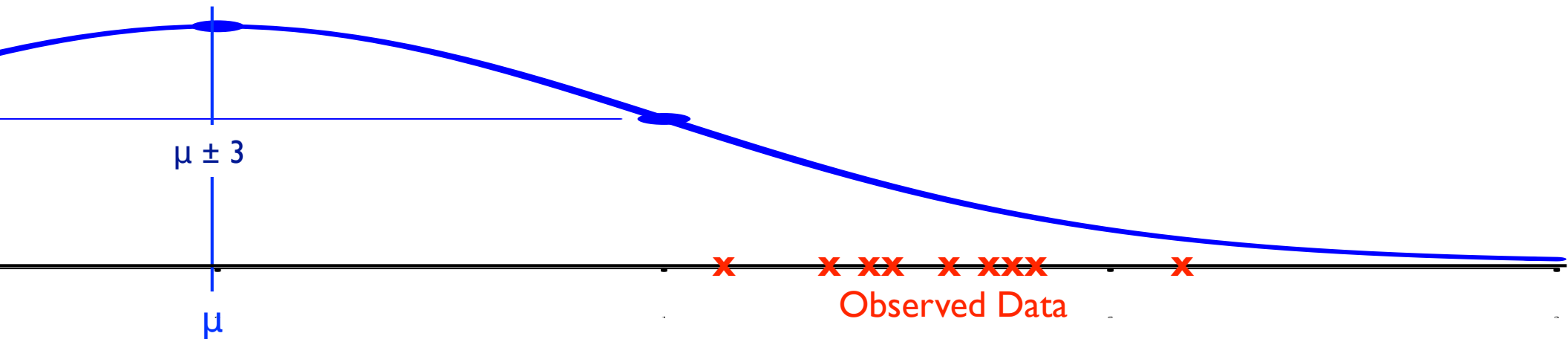
Which is more likely: (a) this?

μ, σ^2 both unknown



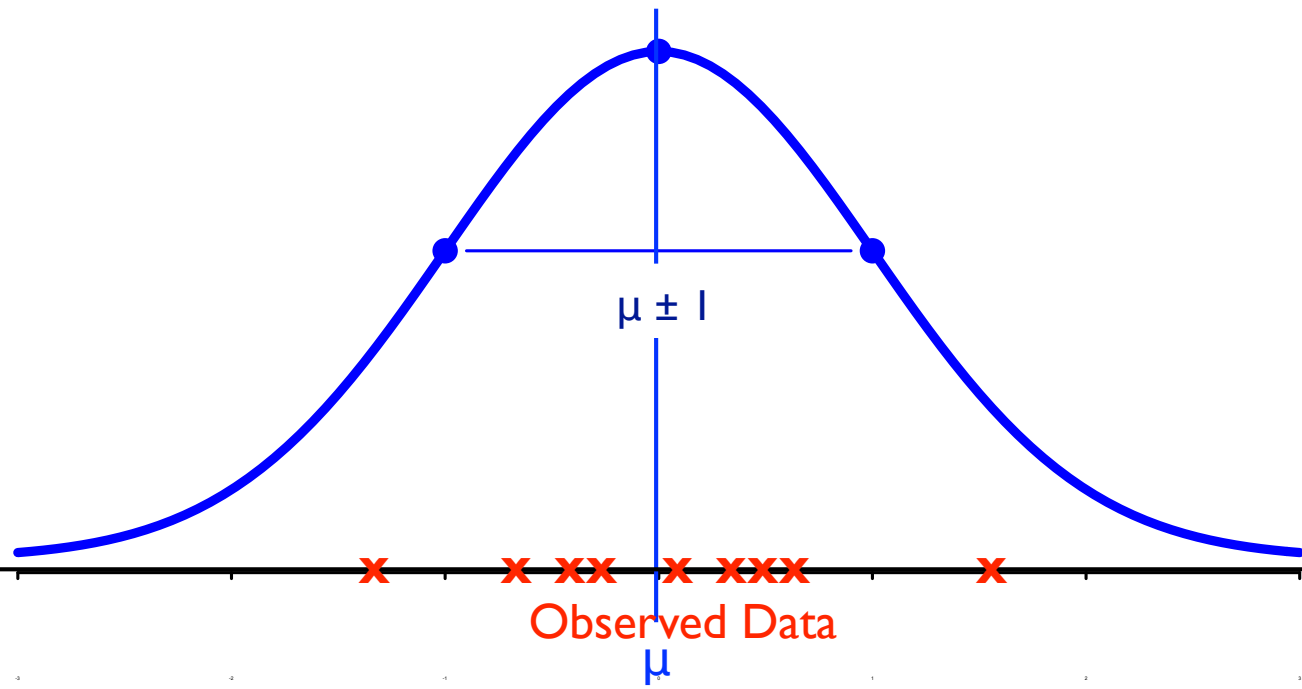
Which is more likely: (b) or this?

μ, σ^2 both unknown



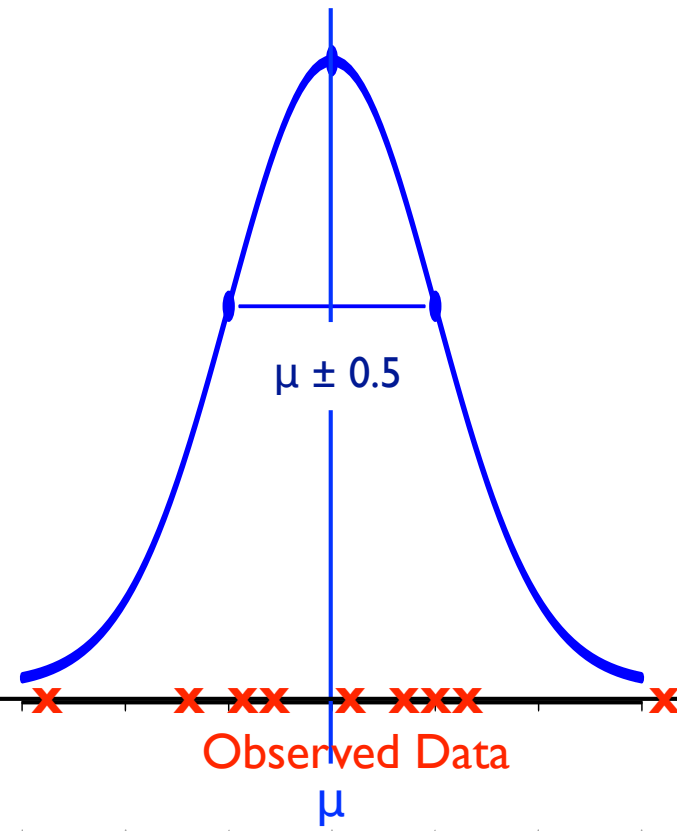
Which is more likely: (c) or this?

μ, σ^2 both unknown



Which is more likely: (d) or *this*?

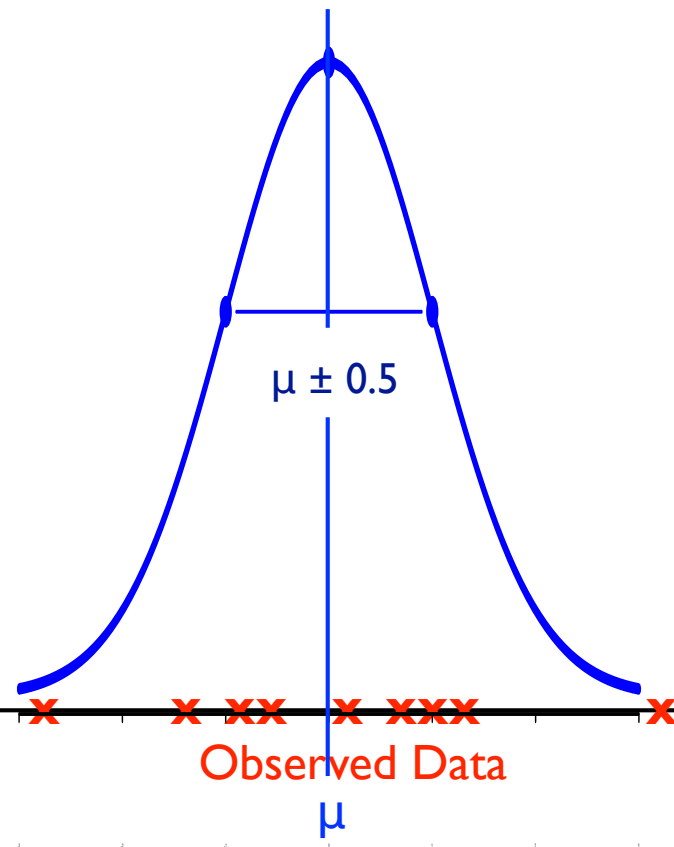
μ, σ^2 both unknown



Which is more likely: (d) or *this*?

μ, σ^2 both unknown

Looks good by eye, but how do I optimize my estimates of μ & $\underline{\underline{\sigma^2}}$?



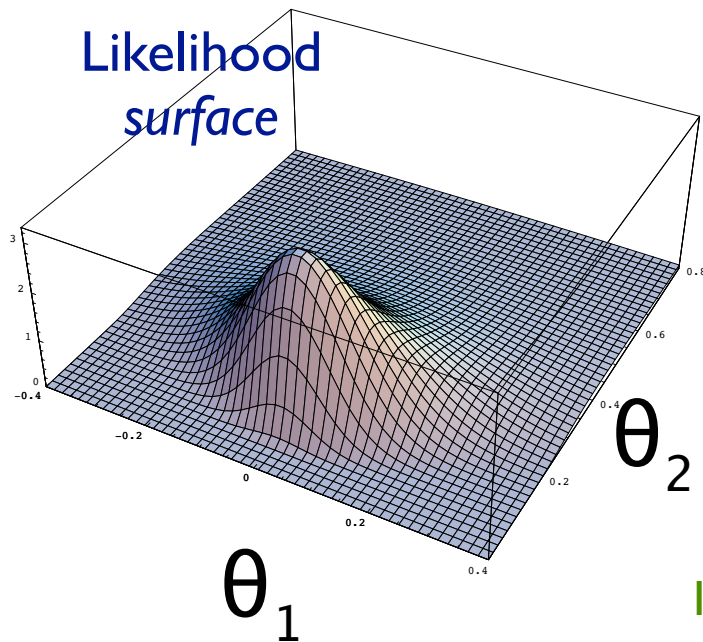
Ex 3: $x_i \sim N(\mu, \sigma^2)$, μ, σ^2 both unknown

$$\ln L(x_1, x_2, \dots, x_n | \theta_1, \theta_2) = \sum_{i=1}^n -\frac{1}{2} \ln(2\pi\theta_2) - \frac{(x_i - \theta_1)^2}{2\theta_2}$$

$$\frac{\partial}{\partial \theta_1} \ln L(x_1, x_2, \dots, x_n | \theta_1, \theta_2) = \sum_{i=1}^n \frac{(x_i - \theta_1)}{\theta_2} = 0$$

$$\hat{\theta}_1 = \left(\sum_{i=1}^n x_i \right) / n = \bar{x}$$

Likelihood
surface



Sample mean is MLE of
population mean, again

In general, a problem like this results in 2 equations in 2 unknowns.
Easy in this case, since θ_2 drops out of the $\partial/\partial\theta_1 = 0$ equation 29

Ex. 3, (cont.)

$$\ln L(x_1, x_2, \dots, x_n | \theta_1, \theta_2) = \sum_{i=1}^n -\frac{1}{2} \ln(2\pi\theta_2) - \frac{(x_i - \theta_1)^2}{2\theta_2}$$

$$\frac{\partial}{\partial \theta_2} \ln L(x_1, x_2, \dots, x_n | \theta_1, \theta_2) = \sum_{i=1}^n -\frac{1}{2} \frac{2\pi}{2\pi\theta_2} + \frac{(x_i - \theta_1)^2}{2\theta_2^2} = 0$$

$$\hat{\theta}_2 = \left(\sum_{i=1}^n (x_i - \hat{\theta}_1)^2 \right) / n = \bar{s}^2$$

*Sample variance is MLE of
population variance*

Ex. 3, (cont.)

$$\ln L(x_1, x_2, \dots, x_n | \theta_1, \theta_2) = \sum_{i=1}^n -\frac{1}{2} \ln(2\pi\theta_2) - \frac{(x_i - \theta_1)^2}{2\theta_2}$$

$$\frac{\partial}{\partial \theta_2} \ln L(x_1, x_2, \dots, x_n | \theta_1, \theta_2) = \sum_{i=1}^n -\frac{1}{2} \frac{2\pi}{2\pi\theta_2} + \frac{(x_i - \theta_1)^2}{2\theta_2^2} = 0$$

$$\hat{\theta}_2 = \left(\sum_{i=1}^n (x_i - \hat{\theta}_1)^2 \right) / n = \bar{s}^2$$

A consistent, but *biased* estimate of population variance.

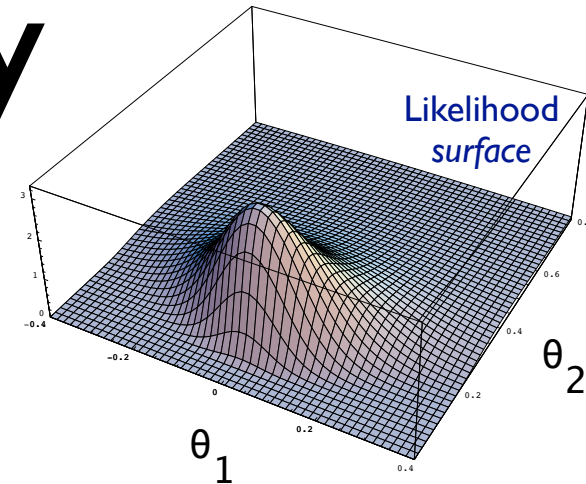
(An example of *overfitting*.) Unbiased estimate is:

$$\hat{\theta}'_2 = \sum_{i=1}^n \frac{(x_i - \hat{\theta}_1)^2}{n - 1}$$

i.e., $\lim_{n \rightarrow \infty}$
= correct

Moral: MLE is a great idea, but not a magic bullet

MLE Summary



MLE is *one* way to estimate *parameters* from *data*

You choose the *form* of the model (normal, binomial, ...)

Math chooses the *value(s)* of parameter(s)

Defining the “Likelihood Function” (based on the pmf or pdf of the model) is often the critical step; the math/algorithms to optimize it are generic

$$\text{Often simply } (d/d\theta)(\log \text{Likelihood}(\text{data}|\theta)) = 0$$

Has the intuitively appealing property that the parameters maximize the *likelihood* of the observed data; basically just assumes your sample is “representative”

Of course, unusual samples will give bad estimates (estimate normal human heights from a sample of NBA stars?) but that is an unlikely event

Often, but not always, MLE has other desirable properties like being *unbiased*, or at least *consistent*

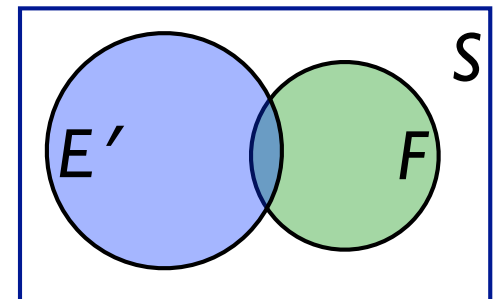
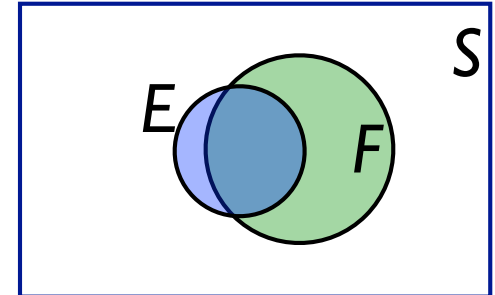
Conditional Probability & Bayes Rule

Conditional probability of E given F: probability that E occurs given that F has occurred.

“Conditioning on F”

Written as $P(E|F)$

Means “P(E has happened, given F observed)”



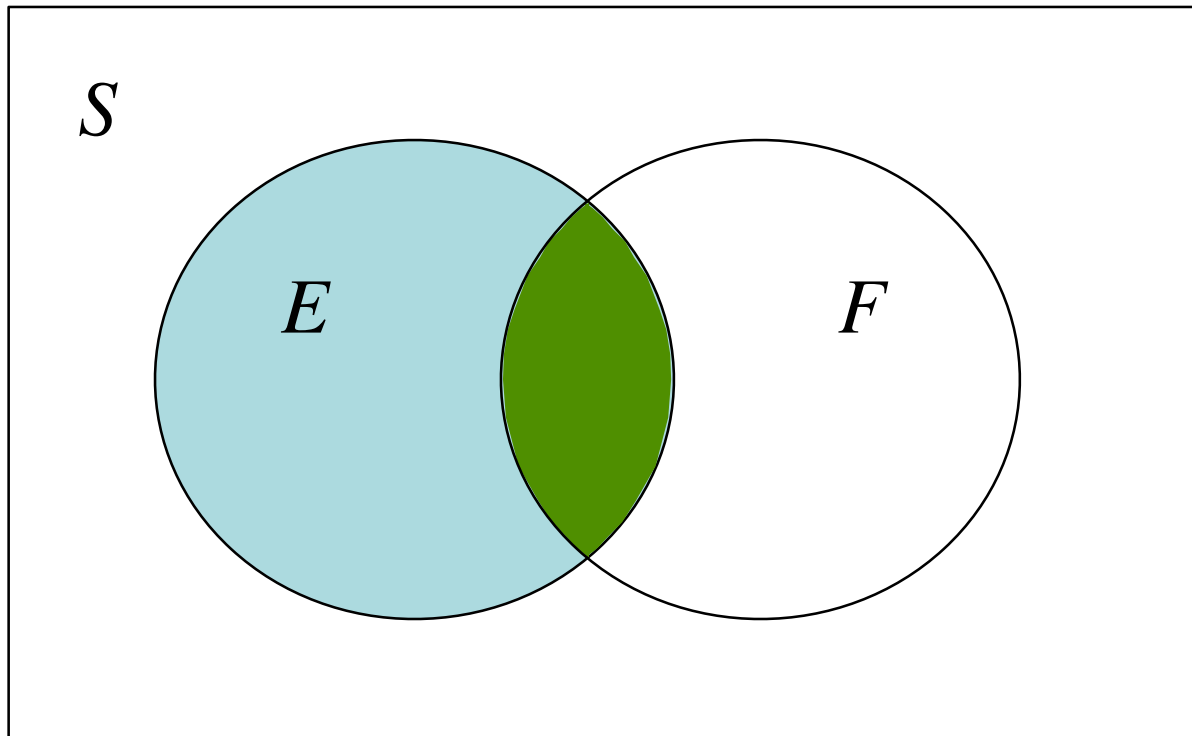
$$P(E | F) = \frac{P(EF)}{P(F)}$$

where $P(F) > 0$

$$P(EF) = P(E | F) P(F)$$

E and F are events in the sample space S

$$E = EF \cup EF^c$$



$$EF \cap EF^c = \emptyset$$

$$\Rightarrow P(E) = P(EF) + P(EF^c)$$

Most common form:

$$P(F | E) = \frac{P(E | F)P(F)}{P(E)}$$

Expanded form (using law of total probability):

$$P(F | E) = \frac{P(E | F)P(F)}{P(E | F)P(F) + P(E | F^c)P(F^c)}$$

Proof:

$$P(F | E) = \frac{P(EF)}{P(E)} = \frac{P(E | F)P(F)}{P(E)}$$

The "EM" Algorithm

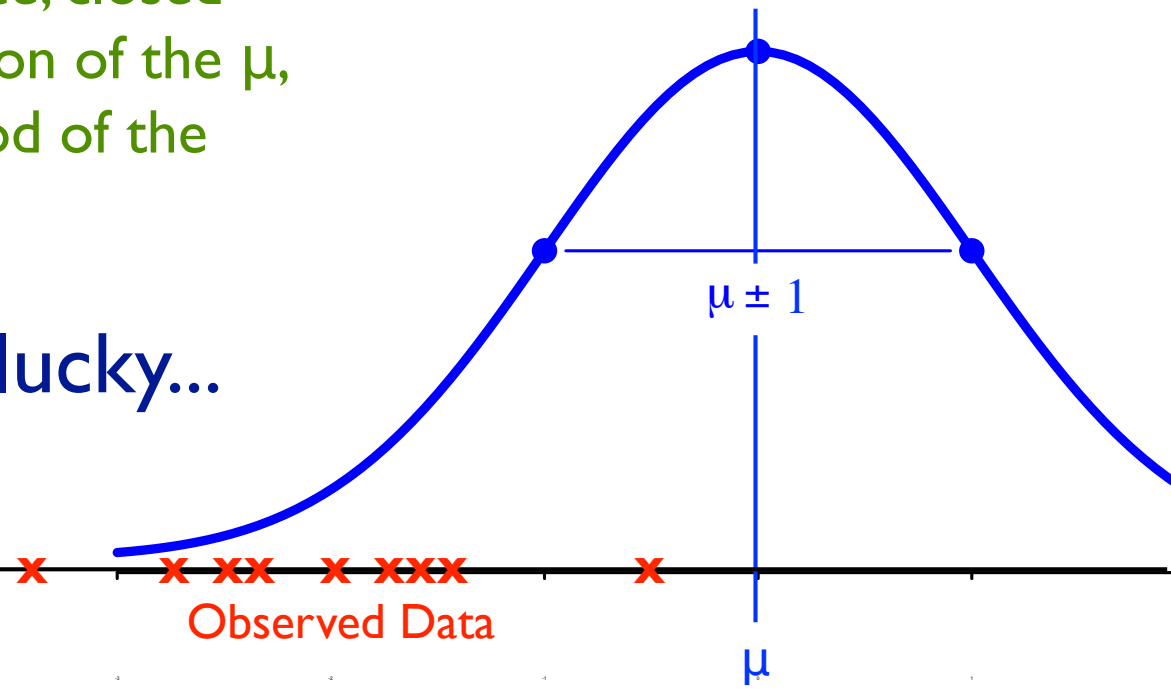
The Expectation-Maximization Algorithm
(for a Two-Component Mixture)

Previously:

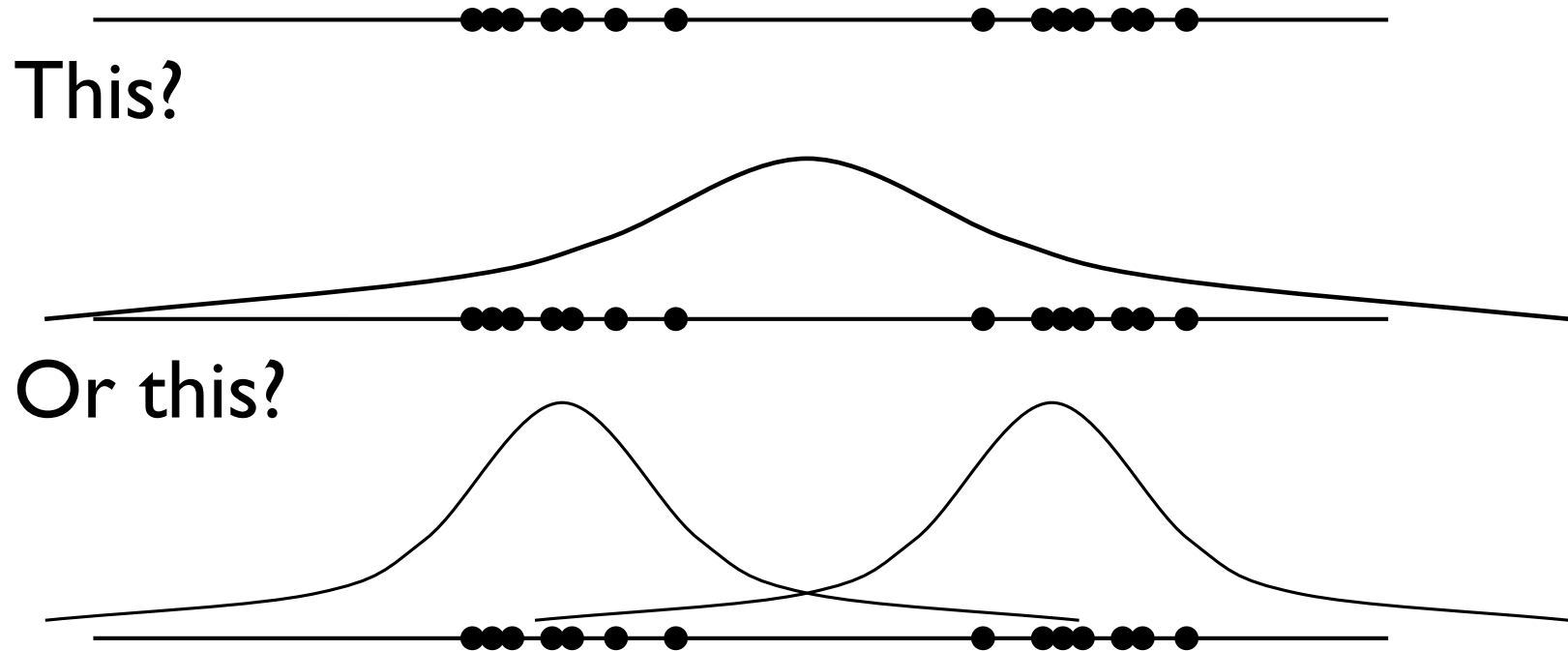
How to estimate μ given data

For this problem, we got a nice, closed form, solution, allowing calculation of the μ , σ that maximize the likelihood of the observed data.

We're not always so lucky...

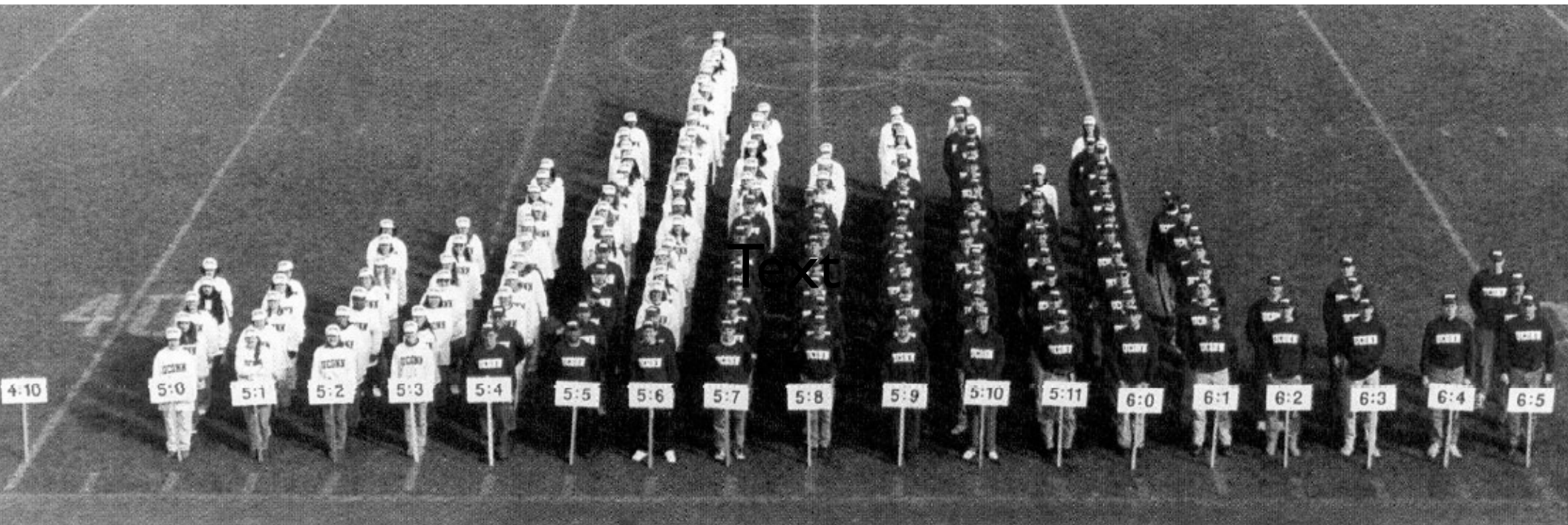


More Complex Example



(A modeling decision, not a math problem...,
but if the later, what math?)

A Living Histogram



male and female genetics students, University of Connecticut in 1996

<http://mindprod.com/jgloss/histogram.html>

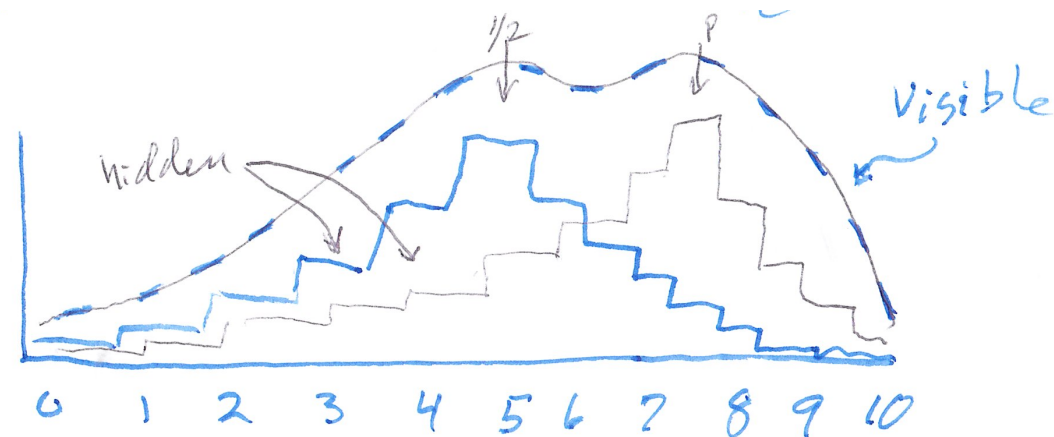
2 Coins: A Binomial Mixture

One fair coin ($p(H)=1/2$), and
one biased coin ($p(H) = p$, fixed but unknown)

For $i = 1, 2, \dots, n$:
pick a coin at random,
flip it 10 times
record $x_i = \#$ of heads

What is MLE for p ?

Expect histogram of
 x_i to look like:



EM as Chicken vs Egg

Hidden Data: let $z_i = 1$ if x_i was from biased coin, else 0

- IF I knew z_i , I could estimate p
(easy: just use x_i s.t. $z_i = 1$)
- IF I knew p , I could estimate z_i
(E.g., if $p = .8$, $x_i \geq 8$ implies z_i more likely 1;
 $x_i \leq 5$ implies z_i more likely 0;
not clear-cut between, but uncertainty is quantifiable.)

Sadly, I know
neither,
... but ...

The "E-M Algorithm": iterate until convergence:

E-step: given (estimated) p , (re)-estimate z 's

M-step: given (estimated) z 's, (re)-estimate p

} Be Optimistic!

The E-Step

$$E = 0 \cdot P(0) + 1 \cdot P(1)$$

↓

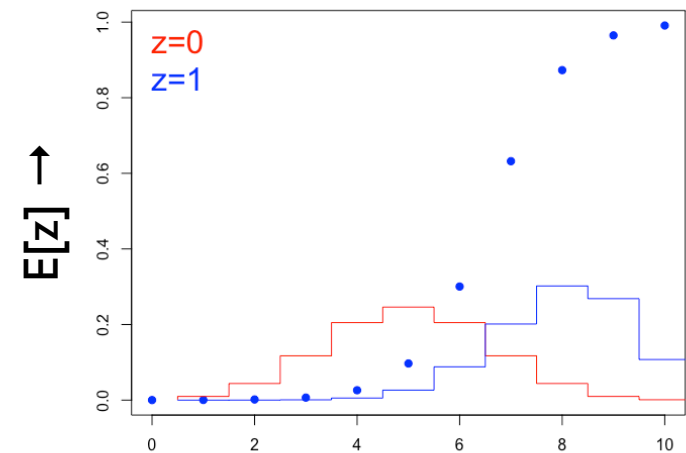
$$E[z_i] = Pr(z_i = 1 | x_i)$$

Bayes

$$= \frac{Pr(x_i | z_i = 1)Pr(z_i = 1)}{Pr(x_i | z_i = 1)Pr(z_i = 1) + Pr(x_i | z_i = 0)Pr(z_i = 0)}$$

$$= \frac{\binom{10}{x_i} \cdot p^{x_i} (1-p)^{10-x_i} \cdot \frac{1}{2}}{\binom{10}{x_i} \cdot p^{x_i} (1-p)^{10-x_i} \cdot \frac{1}{2} + \binom{10}{x_i} \cdot \left(\frac{1}{2}\right)^{10} \cdot \frac{1}{2}}$$

$$= \frac{p^{x_i} (1-p)^{10-x_i}}{p^{x_i} (1-p)^{10-x_i} + 2^{-10}}$$



Math-Hacking the "if "

Let $b(x | p)$ = binomial prob of x heads in 10 flips when $p(H)=p$

As above, $z = 1$ if x was biased, else 0

Then likelihood of x is

$$L(x, z | p) = \text{"if } z == 1 \text{ then } b(x | p) \text{ else } b(x | 1/2)\text{"}$$

Is there a smoother way? Especially, a differentiable way?

Yes! Idea #1:

$$L(x, z | p) = z \cdot b(x | p) + (1-z) \cdot b(x | 1/2)$$

Better still, idea #2:

$$L(x, z | p) = b(x | p)^z \cdot b(x | 1/2)^{(1-z)}$$

equal,
if
 z is 0/1

" θ " here = " p "
on previous slide

The M-Step

$$L(\vec{x}, \vec{z} | \theta) = C \prod_{i=1}^n (\theta^{x_i} (1 - \theta)^{10 - x_i})^{z_i}, \text{ where } C = \prod_{i=1}^n \binom{10}{x_i} \left(\frac{1}{2^{10}}\right)^{1 - z_i}$$

$$\begin{aligned} E[\log L(\vec{x}, \vec{z} | \theta)] &= E \left[\log C + \sum_{i=1}^n z_i (x_i \log \theta + (10 - x_i) \log(1 - \theta)) \right] \\ &= E[\log C] + \sum_{i=1}^n E[z_i] (x_i \log \theta + (10 - x_i) \log(1 - \theta)) \end{aligned}$$

linearity of expectation

$$\frac{d}{d\theta} E[\log L(\vec{x}, \vec{z} | \theta)] = 0 + \sum_{i=1}^n E[z_i] \left(\frac{x_i}{\theta} - \frac{10 - x_i}{1 - \theta} \right)$$

Set to zero and solve, using $E[z_i] = \hat{z}_i$ from E-step. Result (after some algebra):

$$\hat{\theta} = \frac{\sum_{i=1}^n \hat{z}_i \cdot x_i}{\sum_{i=1}^n \hat{z}_i \cdot 10}$$

Intuitively sensible: the estimated fraction of heads from the biased coin is the observed fraction of heads seen overall, after *weighting* by the probability that each observation was indeed from the biased coin.

An E-M "Alg" for This Problem

Input: $x_1, \dots, x_n, 0 \leq x_i \leq 10$

Guess p

Maybe randomly restart
here a few times



repeat until convergence:

E-step: calculate $E[z_1], \dots, E[z_n]$, using latest p

M-step: update p using $E[z_i]$ (the p that maximizes likelihood given those $E[z_i]$)

Alternatively: start by guessing $E[z_i]$, then do M-E-M-...

Suggested exercise(s)

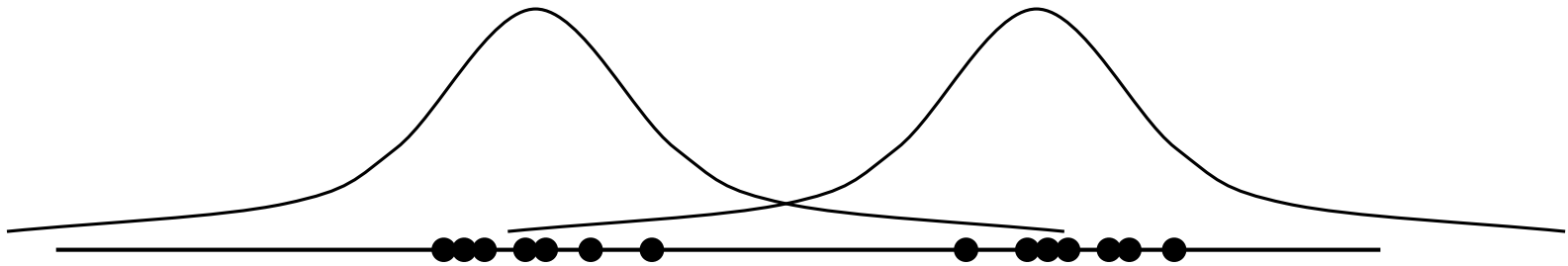
Redo the math assuming *both* coins are biased (but unequally)

Write code to implement either version

Or a spreadsheet, with "fill down" to do a few iterations

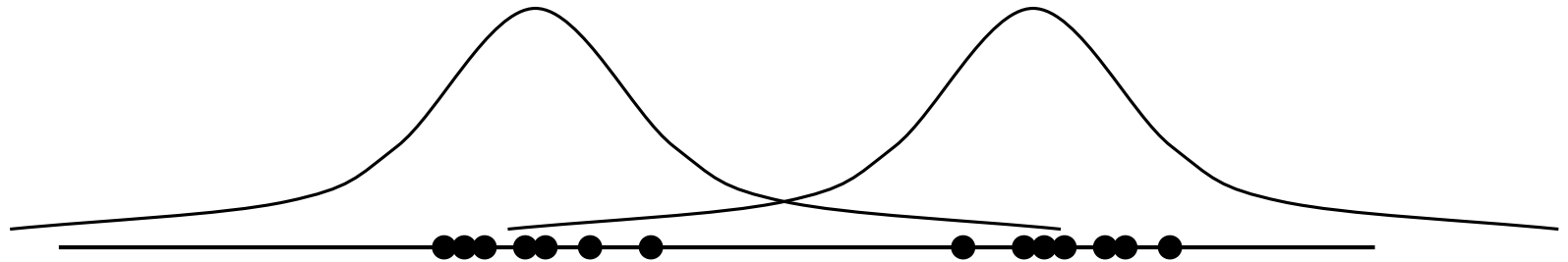
Even in the 1-coin-biased version, there may be multiple local maxima (e.g., consider histogram with a small peak at .25 and large ones at .5 & .8) Does your alg get stuck at local max? How often? Does random restart pragmatically fix this?

EM for a Gaussian Mixture



I have presented the Gaussian mixture example in other courses. I will *NOT* lecture on it in 427, but I'll leave the slides (48-60) here in case you are interested in seeing another example in detail.
Happy to discuss in office hours.

Gaussian Mixture Models / Model-based Clustering



Parameters θ

means	μ_1	μ_2
variances	σ_1^2	σ_2^2
mixing parameters	τ_1	$\tau_2 = 1 - \tau_1$

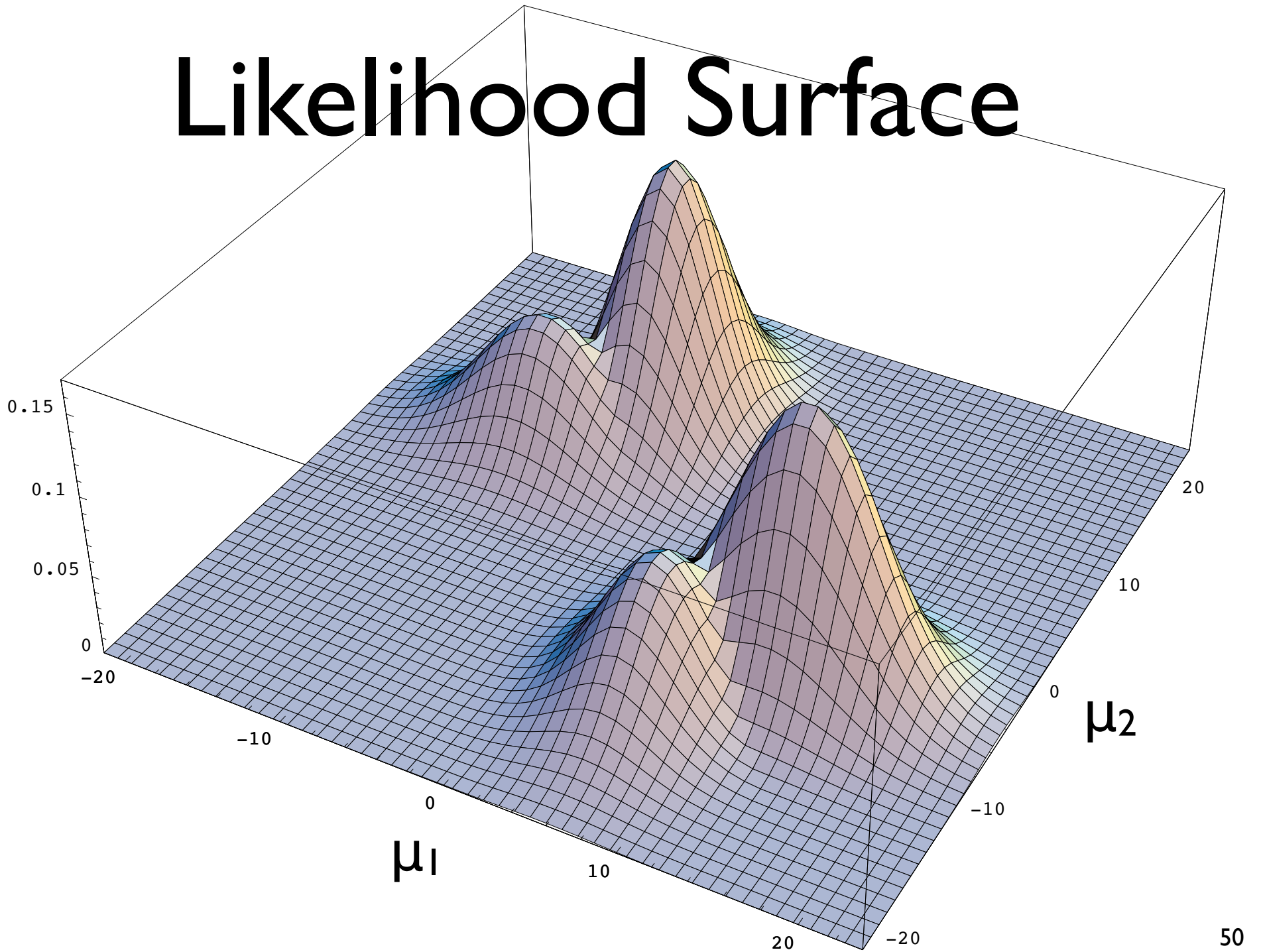
P.D.F. $\xrightarrow{\text{separately}}$ $f(x|\mu_1, \sigma_1^2)$ $f(x|\mu_2, \sigma_2^2)$

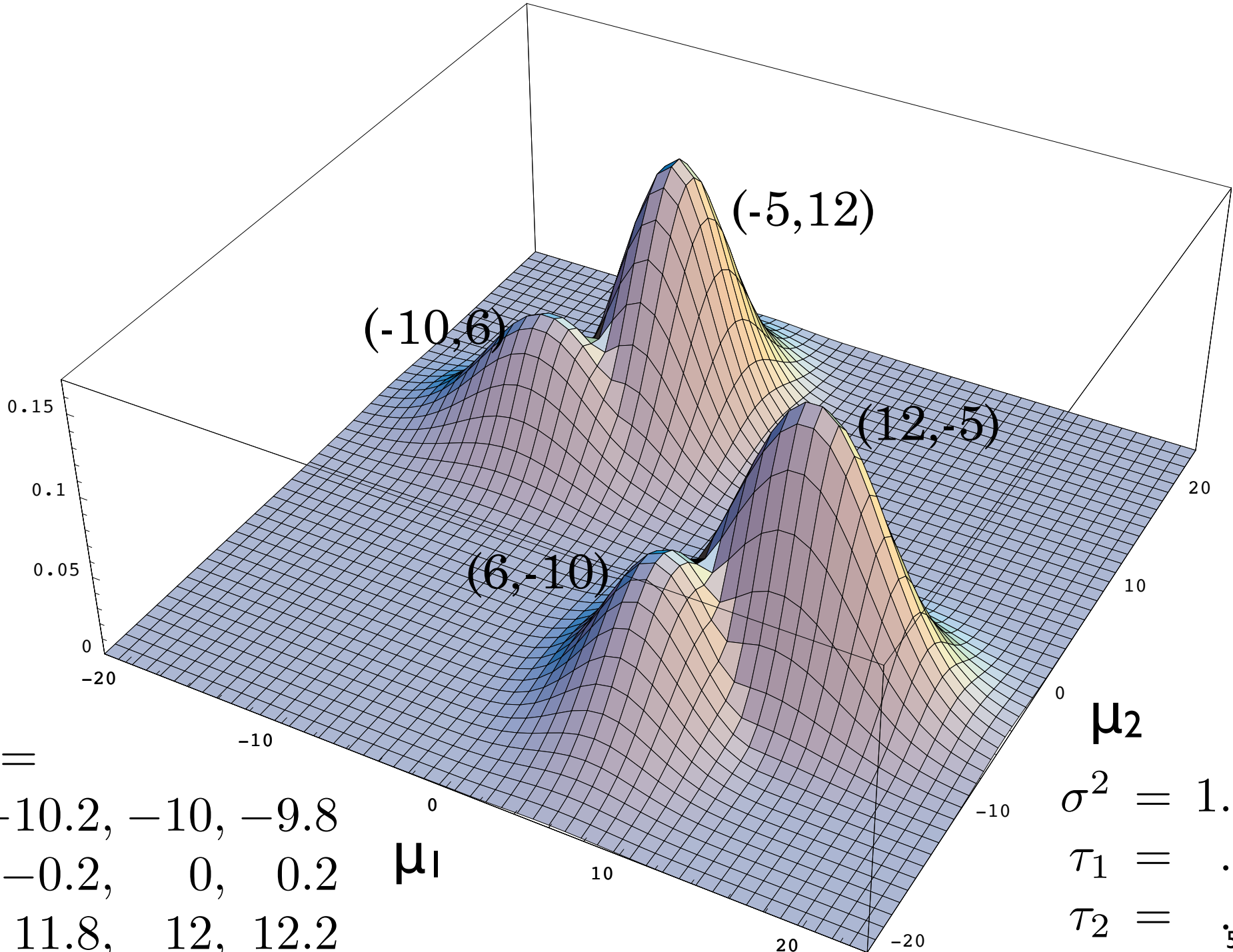
$\xrightarrow{\text{together}}$ $\tau_1 f(x|\mu_1, \sigma_1^2) + \tau_2 f(x|\mu_2, \sigma_2^2)$

Likelihood $\left\{ \begin{aligned} &L(x_1, x_2, \dots, x_n | \mu_1, \mu_2, \sigma_1^2, \sigma_2^2, \tau_1, \tau_2) \\ &= \prod_{i=1}^n \sum_{j=1}^2 \tau_j f(x_i | \mu_j, \sigma_j^2) \end{aligned} \right.$

No closed-form max

Likelihood Surface





$x_i =$
 -10.2, -10, -9.8
 -0.2, 0, 0.2
 11.8, 12, 12.2

μ_1

μ_2
 $\sigma^2 = 1.0$
 $\tau_1 = .5$
 $\tau_2 = .5$

A What-If Puzzle

Likelihood

$$L(x_1, x_2, \dots, x_n | \overbrace{\mu_1, \mu_2, \sigma_1^2, \sigma_2^2, \tau_1, \tau_2}^{\theta})$$
$$= \prod_{i=1}^n \sum_{j=1}^2 \tau_j f(x_i | \mu_j, \sigma_j^2)$$

Messy: no closed form solution known for finding θ maximizing L

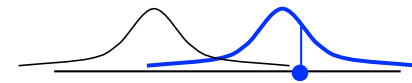
But *what if* we knew the *hidden data*?

$$z_{ij} = \begin{cases} 1 & \text{if } x_i \text{ drawn from } f_j \\ 0 & \text{otherwise} \end{cases}$$

EM as Egg vs Chicken

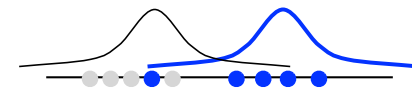
IF parameters θ known, could estimate z_{ij}

E.g., $|x_i - \mu_1|/\sigma_1 \gg |x_i - \mu_2|/\sigma_2 \Rightarrow P[z_{i1}=1] \ll P[z_{i2}=1]$



IF z_{ij} known, could estimate parameters θ

E.g., only points in cluster 2 influence μ_2, σ_2



But we know neither; (optimistically!) iterate:

E-step: calculate expected z_{ij} , given parameters

M-step: calculate “MLE” of parameters, given $E(z_{ij})$

Overall, a clever “hill-climbing” strategy

Not “EM,” but may help clarify concepts

Simple Version: “Classification EM”

If $E[z_{ij}] < .5$, pretend $z_{ij} = 0$; $E[z_{ij}] > .5$, pretend it's 1

I.e., *classify* points as component 1 or 2

Now recalc θ , assuming that partition (standard MLE)

Then recalc $E[z_{ij}]$, assuming that θ

Then re-recalc θ , assuming new $E[z_{ij}]$, etc., etc.

“K-means clustering,” essentially

“Full EM” is slightly more involved, (to account for uncertainty in classification) but this is the crux.

Another contrast: HMM parameter estimation via “Viterbi” vs “Baum-Welch” training. In both, “hidden data” is “which state was it in at each step?” Viterbi is like E-step in classification EM: it makes a single state prediction. B-W is full EM: it captures the uncertainty in state prediction, too. For either, M-step maximizes HMM emission/transition probabilities, assuming those fixed states (Viterbi) / uncertain states (B-W).

Full EM

x_i 's are known; θ unknown. Goal is to find MLE θ of:

$$L(x_1, \dots, x_n \mid \theta) \quad \text{(hidden data likelihood)}$$

Would be easy *if* z_{ij} 's were known, i.e., consider:

$$L(x_1, \dots, x_n, z_{11}, z_{12}, \dots, z_{n2} \mid \theta) \quad \text{(complete data likelihood)}$$

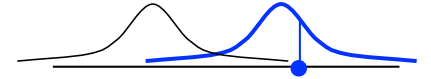
But z_{ij} 's aren't known.

Instead, maximize *expected* likelihood of visible data

$$E(L(x_1, \dots, x_n, z_{11}, z_{12}, \dots, z_{n2} \mid \theta)),$$

where expectation is over distribution of hidden data (z_{ij} 's)

i.e., average over possible, but hidden z_{ij} 's



The E-step:

Find $E(z_{ij})$, i.e., $P(z_{ij}=1)$

Assume θ known & fixed

A (B): the event that x_i was drawn from f_1 (f_2)

D: the observed datum x_i

Expected value of z_{i1} is $P(A|D)$

$$E = 0 \cdot P(0) + 1 \cdot P(1)$$

$$E[z_{i1}] = P(A|D) = \frac{P(D|A)P(A)}{P(D)}$$

$$\begin{aligned} P(D) &= P(D|A)P(A) + P(D|B)P(B) \\ &= f_1(x_i|\theta_1)\tau_1 + f_2(x_i|\theta_2)\tau_2 \end{aligned}$$

Repeat
for
each
 $z_{i,j}$

Note: denominator = sum of numerators – i.e. that which normalizes sum to 1 (typical Bayes)

Complete Data Likelihood

Recall:

$$z_{1j} = \begin{cases} 1 & \text{if } x_1 \text{ drawn from } f_j \\ 0 & \text{otherwise} \end{cases}$$

so, correspondingly,

$$L(x_1, z_{1j} | \theta) = \begin{cases} \tau_1 f_1(x_1 | \theta) & \text{if } z_{11} = 1 \\ \tau_2 f_2(x_1 | \theta) & \text{otherwise} \end{cases}$$

equal, if z_{ij} are 0/1



Formulas with “if’s” are messy; can we blend more smoothly?

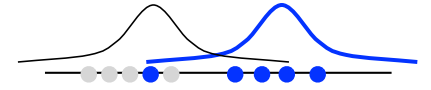
Yes, many possibilities. Idea 1:

$$L(x_1, z_{1j} | \theta) = z_{11} \cdot \tau_1 f_1(x_1 | \theta) + z_{12} \cdot \tau_2 f_2(x_1 | \theta)$$

Idea 2 (Better):

$$L(x_1, z_{1j} | \theta) = (\tau_1 f_1(x_1 | \theta))^{z_{11}} \cdot (\tau_2 f_2(x_1 | \theta))^{z_{12}}$$

M-step:



Find θ maximizing $E(\log(\text{Likelihood}))$

(For simplicity, assume $\sigma_1 = \sigma_2 = \sigma; \tau_1 = \tau_2 = \tau = 0.5$)

$$L(\vec{x}, \vec{z} | \theta) = \prod_{i=1}^n \left(\frac{\tau}{\sqrt{2\pi\sigma^2}} \exp \left(- \sum_{j=1}^2 z_{ij} \frac{(x_i - \mu_j)^2}{2\sigma^2} \right) \right)$$

$$E[\log L(\vec{x}, \vec{z} | \theta)] = E \left[\sum_{i=1}^n \left(\log \tau - \frac{1}{2} \log(2\pi\sigma^2) - \sum_{j=1}^2 z_{ij} \frac{(x_i - \mu_j)^2}{2\sigma^2} \right) \right]$$

wrt dist of z_{ij}

$$= \sum_{i=1}^n \left(\log \tau - \frac{1}{2} \log(2\pi\sigma^2) - \sum_{j=1}^2 E[z_{ij}] \frac{(x_i - \mu_j)^2}{2\sigma^2} \right)$$

Find θ maximizing this as before, using $E[z_{ij}]$ found in E-step. Result:

$$\mu_j = \frac{\sum_{i=1}^n E[z_{ij}] x_i}{\sum_{i=1}^n E[z_{ij}]} \quad (\text{intuit: avg, weighted by subpop prob})$$

M-step: calculating mu's

$$\mu_j = \sum_{i=1}^n E[z_{ij}]x_i / \sum_{i=1}^n E[z_{ij}]$$

In words: μ_j is the average of the observed x_i 's, weighted by the probability that x_i was sampled from component j .

old E's

							row sum	avg
E[z _{i1}]	0.99	0.98	0.7	0.2	0.03	0.01	2.91	
E[z _{i2}]	0.01	0.02	0.3	0.8	0.97	0.99	3.09	
x _i	9	10	11	19	20	21	90	15
E[z _{i1}]x _i	8.9	9.8	7.7	3.8	0.6	0.2	31.02	10.66
E[z _{i2}]x _i	0.1	0.2	3.3	15.2	19.4	20.8	58.98	19.09

new μ's

2 Component Mixture

$$\sigma_1 = \sigma_2 = 1; \tau = 0.5$$

		mu1	-20.00		-6.00		-5.00		-4.99
		mu2	6.00		0.00		3.75		3.75
x1	-6	z11		5.11E-12		1.00E+00		1.00E+00	
x2	-5	z21		2.61E-23		1.00E+00		1.00E+00	
x3	-4	z31		1.33E-34		9.98E-01		1.00E+00	
x4	0	z41		9.09E-80		1.52E-08		4.11E-03	
x5	4	z51		6.19E-125		5.75E-19		2.64E-18	
x6	5	z61		3.16E-136		1.43E-21		4.20E-22	
x7	6	z71		1.62E-147		3.53E-24		6.69E-26	

Essentially converged in 2 iterations

⇒⇒ (Excel spreadsheet on course web)

EM Summary

Fundamentally, maximum likelihood parameter estimation; broader than just these examples

Useful if 0/1 hidden data, and if analysis would be more tractable if 0/1 hidden data z were known

Iterate:

E-step: estimate $E(z)$ for each z , given θ

M-step: estimate θ maximizing $E[\log \text{likelihood}]$

given $E[z]$ [where “ $E[\log L]$ ” is wrt random $z \sim E[z] = p(z=1)$]

Bayes

MLE

EM Issues

Under mild assumptions (e.g., DEKM sect 11.6), EM is guaranteed to increase likelihood with every E-M iteration, hence will *converge*.

But it may converge to a *local*, not global, max.
(Recall the 4-bump surface...)

Issue is intrinsic (probably), since EM is often applied to *NP-hard* problems (including clustering, above and motif-discovery, soon)

Nevertheless, widely used, often effective, esp. with *random restarts*