Review: Conditional Probability:

P[we see an elephant today| we are in the Serengeti]
P[we see an elephant today| we are in Manhattan]

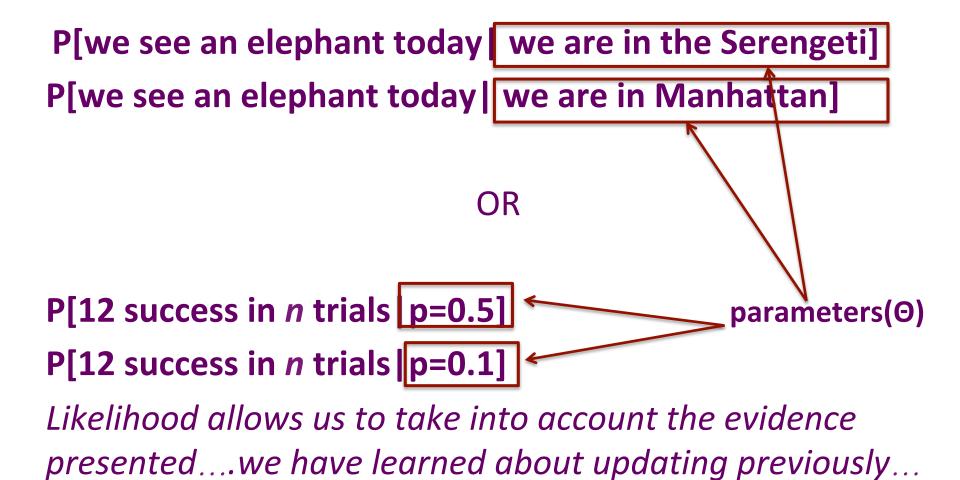
OR

Back to the coin toss world: if our coin was fair, i.e. P(heads) = 0.5 or unfair, i.e. P(heads) = 0.1

P[12 success in n trials | p=0.5] P[12 success in n trials | p=0.1]

Review: Conditional Probability:

The best parameter value is the one that fits the data the best



- Conditional probability
- Components of Bayes':

$$P[A|B] = \underline{P[B|A]P[A]}$$
$$P[B]$$

If A = parameter set, O

And \mathbf{B} = observed data, \mathbf{y}

Then we can re-write the above formula as:

$$P[\Theta|y] = \underline{P[y|\Theta]P[\Theta]}$$
$$P[y]$$

- Conditional probability
- Components of Bayes': Likelihood

$$P[A|B] = P[B|A]P[A]$$
 Prior $P[B]$

If A = parameter set, Θ

Data

And **B** = observed data, **y**

Then we can re-write the above formula as:

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Where \mathbf{A} = parameter set, $\mathbf{\Theta}$ and \mathbf{B} = observed data, \mathbf{y} Then we can re-write the above formula as:

$$P[\Theta|y] = \underline{P[y|\Theta]P[\Theta]}$$
$$P[y]$$

Re-writing hypothesis parameters and data:

$$P[H_i|Data] = P[H_i|and|Data] = P[Data|H_i]P[H_i]$$

 $P[Data]$ $P[Data]$

You can use this to test ratios of likelihood:

$$\frac{P[H_1|Data]}{P[H_2|Data]} = \frac{P[Data|H_1]}{P[H_2]} \frac{P[H_1]}{P[H_2]}$$
Prior

(sites on a sequence, for instance):

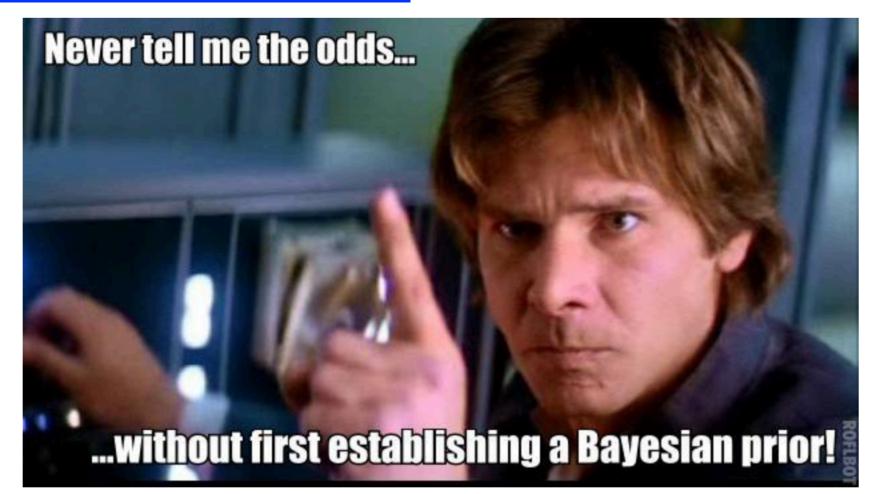
dominates so prior

doesn't influence regults

"Never tell me the odds!"

Star Wars fans:

http://www.countbayesie.com/blog/2015/2/18/hans-solo-and-bayesian-priors



 Quantifies data support for a particular value of a parameter (considers many possible hypotheses, not just one)

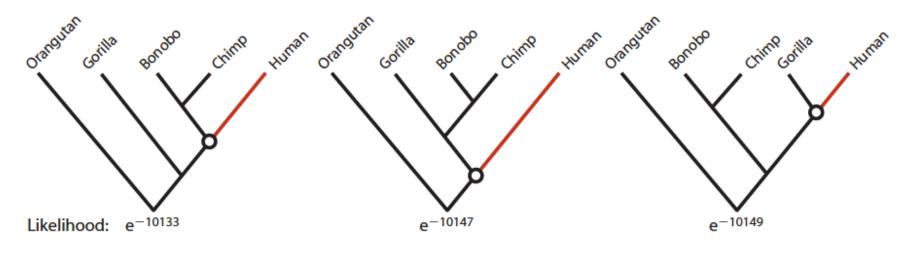
- Probability of obtaining observed data if parameter(s)
 were equal to that specific value
 - Compare to alternate values of the parameter
 - High for values close to the true population parameter and low for values far from it
 - Maximum likelihood estimate: best estimate for parameter

$$L(\theta, data) = L(\theta | data) = P(data | \theta = \theta_0)$$

Does not assume normally distributed data Major Applications:

1. Phylogeny estimation

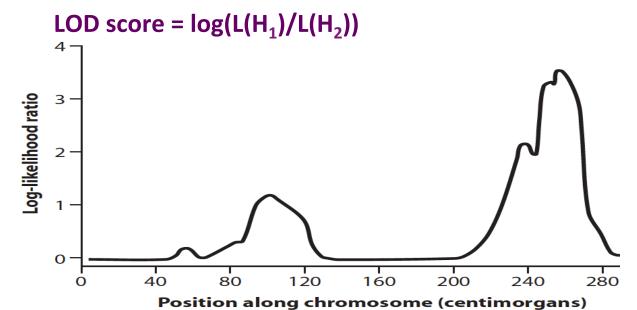
L(Tree=i|Sequence Data|Tree=i)



Major Applications:

2. Gene mapping

- "markers" differ between people with disease and those without
- Comparing two hypotheses:
 - H₁: disease gene is present at a particular site
 - H₂: no disease gene is present at that site
 - Review frequency data in a sliding window along the genome
 - LOD score: log of the ratio of these two hypotheses



Major Applications:

2. Gene mapping

- H₁: disease gene is present at a particular site
- H₂: no disease gene is present at that site
- Review frequency data in a sliding window along the genome
- LOD score: log of the ratio of these two hypotheses

LOD score =
$$log(L(H_1)/L(H_2))$$

This can be thought of as an 'odds ratio' in favour of hypothesis 1 over hypothesis 2

$$P(H_1|D) = P(D|H_1) P(H_1)$$

 $P(H_2|D) P(D|H_2) P(H_2)$

- All statistical estimates that we have encountered so far are maximum likelihood estimates
 - Excellent properties such as consistency and precision as data → ∞

Requires a probability model

- Ex. Mutation models in DNA sequence changes
- Ex. Binomial distribution for proportions
- Determines the parameter value that yields the largest L.
 - A particular data set supports one hypothesis better than another if the likelihood of that hypothesis is higher than the likelihood of the other hypothesis

Very versatile

Example: 11 coin tosses with a supposedly fair coin **HHTTHTHHTTT**

What is the likelihood of this sequence?

Assume coin tosses are independent.

Example: 11 coin tosses with a supposedly fair coin

HHTTHTHHTTT

What is the likelihood of this sequence?

Assume coin tosses are independent.

L(p=0.5|HHTTHTHHTTT) = P(HHTTHTHHTTT|p=0.5)
= pp(1-p)(1-p)p(1-p)pp(1-p)(1-p)(1-p)
=
$$p^{5}(1-p)^{6}$$

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To determine Max. likelihood you are going to need to use some simple **calculus**...followed by some **algebra**

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1.
$$\underline{dL} = 5p^4(1-p)^6 - 6p^5(1-p)^5$$
 dp

2.
$$\underline{dL} = 5p^4(1-p)^6 - 6p^5(1-p)^5 = 0$$
 dp

$$= p^4(1-p)^5[5(1-p)-6p] = 0$$

$$\hat{p} = \frac{5}{11}$$

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$$ln(L)=5lnp+6ln(1-p)$$

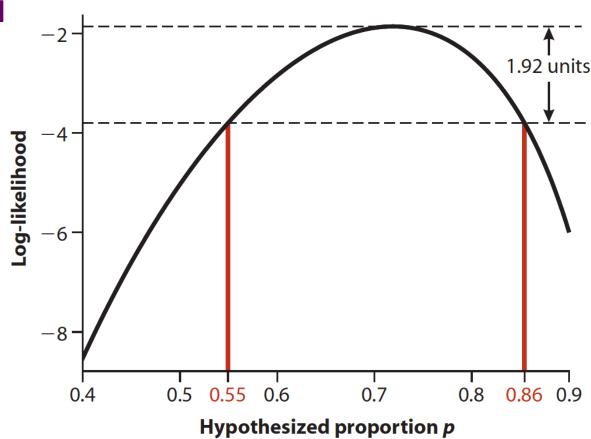
 $d(lnL) = 5 - 6 = 0$
 $p (1-p)$

$$\hat{p} = \frac{5}{11}$$

- Log likelihood is computationally easier
 - Turns multiplication into addition
- Log-likelihood curve
 - Allows us to calculate interval estimate: likelihood-based

confidence interval

• Use $X_{1,\alpha}^2/2$



Example: Counting elephants via dung 'genetic fingerprints'

- originally identify 27 elephants
- sample 74, 15 of which were already sampled in original group of 27 ('resampled')

1. Probability model

- Random sampling
- Population is constant
- Recapture number is 15 out of 74 in our sample
- what we want to know: likelihood of true N is the probability of obtaining this proportion of recaptures for different N values
- This is the difficult step
- Likelihood
- Maximum Likelihood estimate

Example: Counting elephants via dung 'genetic fingerprints'

- originally identify 27 elephants
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- 1. Probability model
 - Recapture number is 15 out of 74 in our sample
 - what we want to know to determine the true N is the probability of obtaining this proportion of recaptures for different N values
 - This is the difficult step; in this case already done for us!

$$P[Y \mid N] = L[N \mid Y] = \frac{\binom{n_1}{Y} \binom{N - n_1}{n_2 - Y}}{\binom{N}{n_2}}$$

- 2. Likelihood
- Maximum Likelihood estimate

Example: Counting elephants via dung 'genetic fingerprints'

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- sample 74, 15 of which were already sampled in original group of 27 ('resampled')
- 1. Probability model
 - Recapture number is 15 out of 74 in our sample
 - what we want to know to determine the true N is the probability of obtaining this proportion of recaptures for different N values
- 2. (log) Likelihood

$$\ln L[N \mid Y] = \ln {\binom{n_1}{Y}} + \ln {\binom{N - n_1}{n_2 - Y}} - \ln {\binom{N}{n_2}}$$

$$\ln L[N \mid Y] = \ln {\binom{27}{15}} + \ln {\binom{N - 27}{74 - 27}} - \ln {\binom{N}{74}}$$

3. Maximum Likelihood estimate

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- originally identify 27 elephants
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- 1. Probability model
 - Recapture number is 15 out of 74 in our sample
 - what we want to know to determine the true N is the probability of obtaining this proportion of recaptures for different N values
- 2. Likelihood

$$ln L[N \mid Y] = ln \begin{bmatrix} 27 \\ 15 \end{bmatrix} + ln \begin{bmatrix} N-27 \\ 74-27 \end{bmatrix} - ln \begin{bmatrix} N \\ 74 \end{bmatrix}$$

3. Maximum Likelihood estimate

$$\hat{N} = 133$$