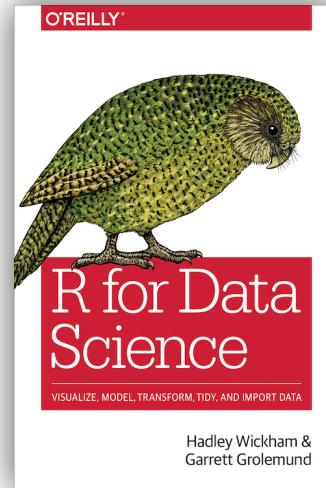
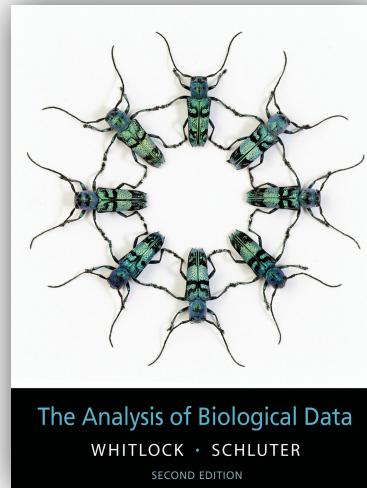


Data Science in Bioinformatics

Palle Villesen & Thomas Bataillon

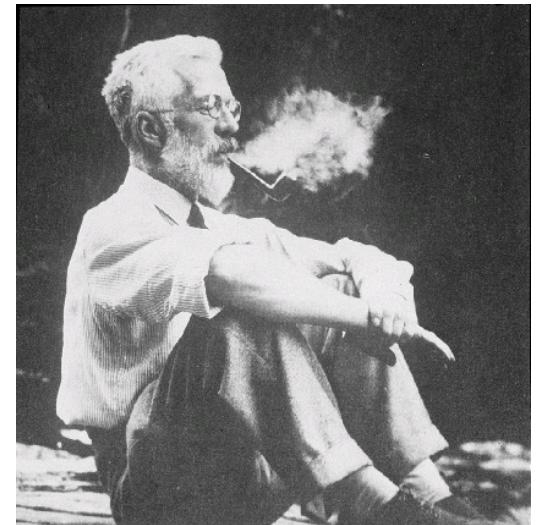


Outline for week 13

- Tuesday: Chapter 20 Likelihood
 - Wasp example (R code)
 - Human trios (R markdown)
 - Important generalizations
- Thursday session is
 - Exercises on likelihood
 - Final assignment (prepare / post Qs)
- We need your feedback

What is likelihood ? Why Bother?

- A general framework for **parameter estimation** and **hypothesis testing**
- An “old” idea (R.A. Fisher 1920s) that went a long way...
- Why likelihood is your “friend”
 - Think clearly about your data
 - Efficient way of extracting information from the data
 - Likelihood is often “hidden” behind tools you use...



Likelihood in a nutshell

- Choose a model M_θ for your data D
- Write down the probability of your data $P(D)$
- Figure out which parameter(s) value(s) of M_θ maximize $P(D)$ (= make the data most likely)
- Distrust your model !
- “Wash, rinse and repeat ...”

Which *model* behinds these data ?

=

Which probability distributions ?

- Height measurements of different individuals from a single population
 - Allele frequencies in a sample taken from a (large) population
 - Sex ratio in a progeny
 - Presence/ absence of a species in a locality
-

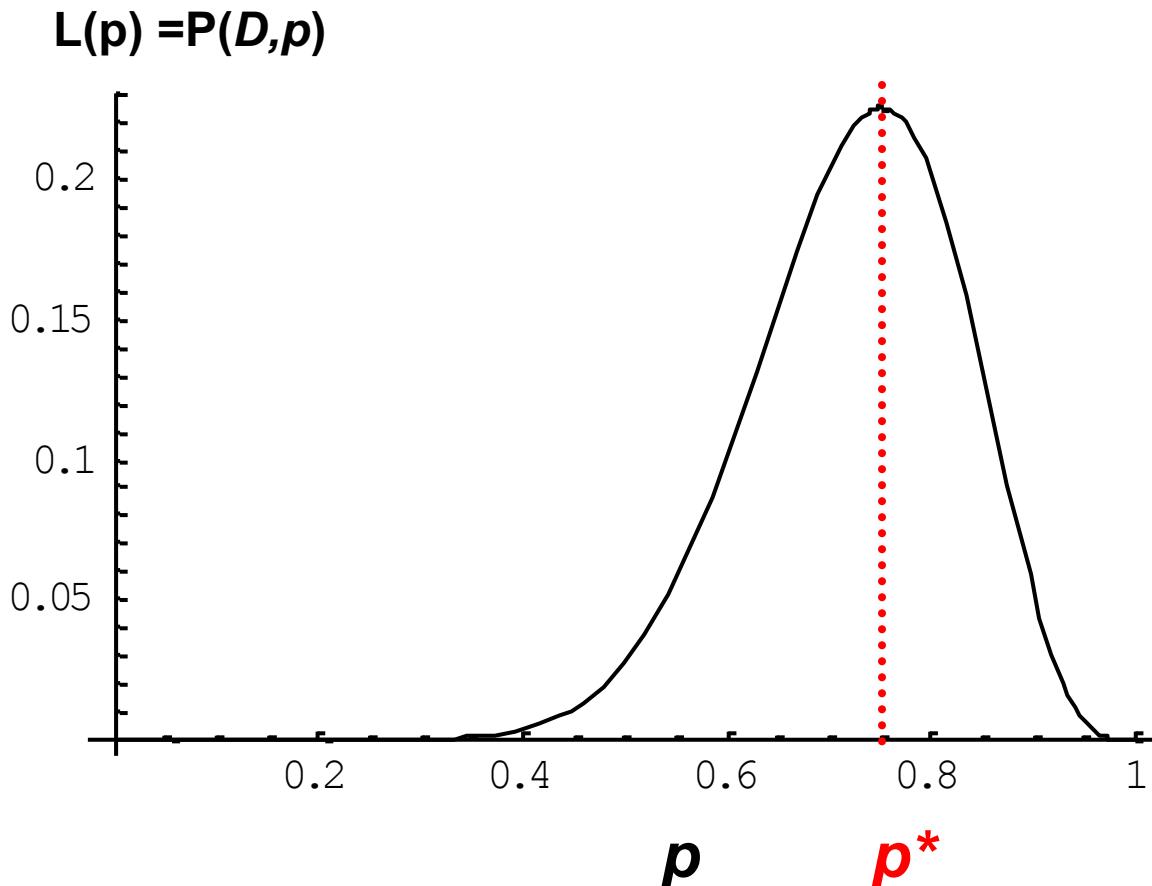
The Binomial distribution

- “Thought Experiment”:
 - n independent trials with probability of success p for each trial
 - X is a **random variable**
 - Formally $X \sim B(n,p)$
- What do we know about X ?
 - $E(X) = n p$ $V(X) = n p (1-p)$
 - $P(X=i) = n!/i!(n-i)! p^i (1-p)^{n-i}$
 - if n is large (np CONSTANT) , X becomes Poisson or even “normal”

The Likelihood principle

- $L = P(D; \theta)$ is a function of data D and parameters θ
- Likelihood principle:
 - The Data D is fixed
 - Choose the parameter(s) value(s) θ^* that make the data D most likely--> Maximize L (THAT's IT !)
- The **invariance principle**: if alpha is a parameter and beta=f(alpha), with f continuous and monotonic, then
 $ML_{\text{beta}} = f(ML_{\text{alpha}})$

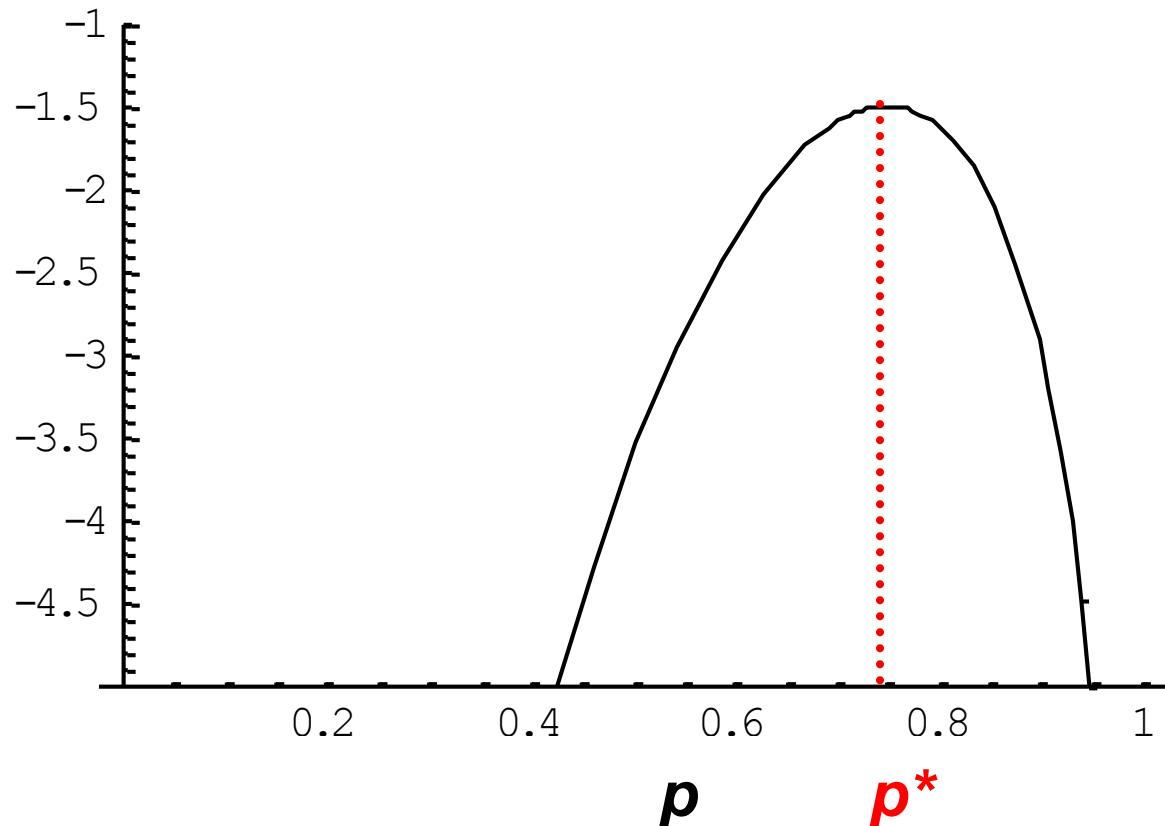
Visualizing the likelihood function



NB: L is a **continuous** function defined for p in $[0,1]$

Visualizing the (natural) log Likelihood function

$\text{LogL}=\text{Log}[P(D)]$

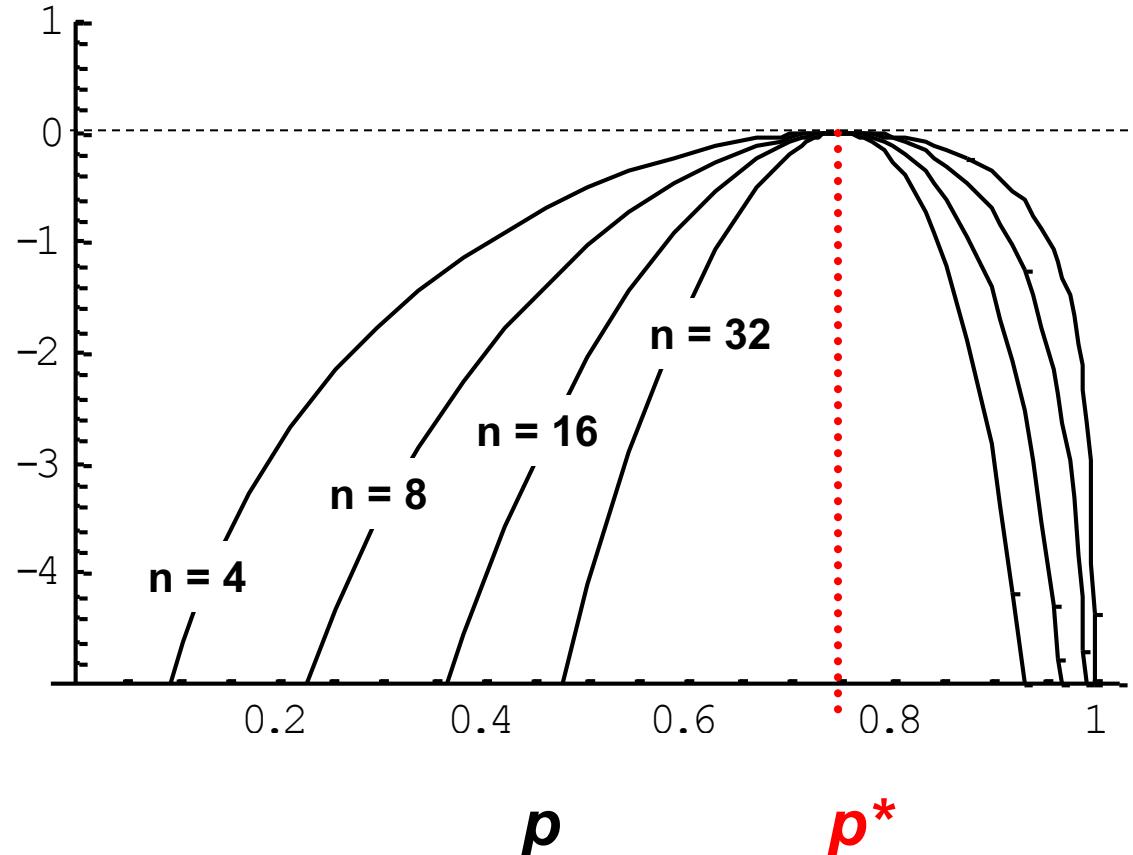


Maximizing the likelihood

- Binomial example (continued)
- Finding the maximum of a function
 1. set the derivative to zero,
 2. check for a maximum.
- Generalization MLEs for several variables...

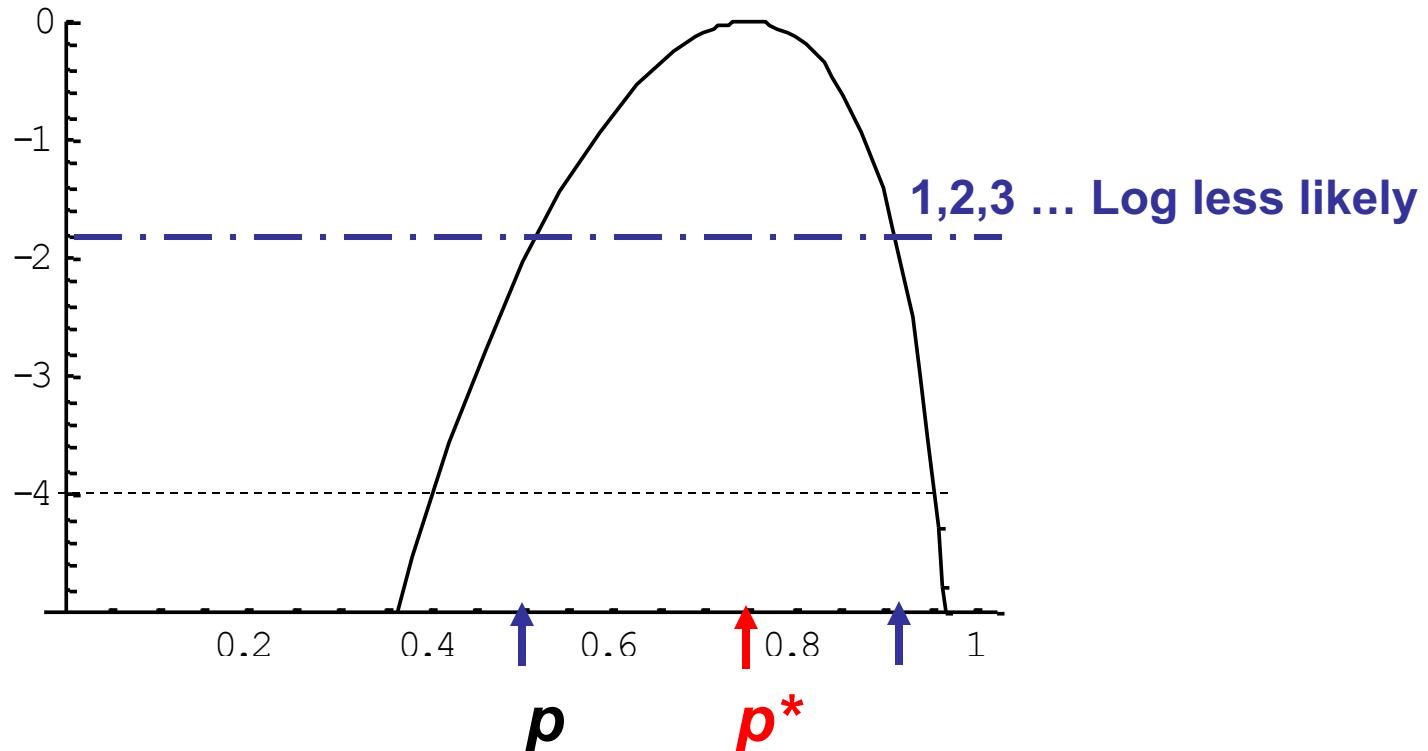
The curvature of Likelihood function reflects the amount of info in your data

$\text{Log}[L(p)] - \text{Log}[L(p^*)]$



Visualizing the log Likelihood function rescaled

$\text{Log}[L(p)] - \text{Log}[L(p^*)]$

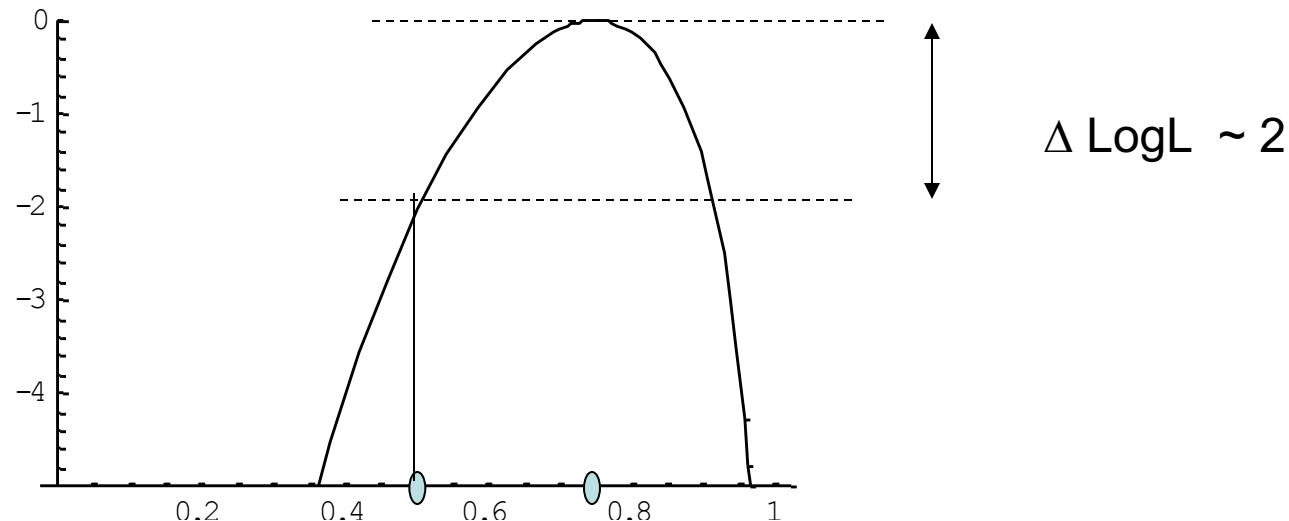


The likelihood profile (graph)

- Intuition: curvature says something about the precision on ML estimates
- General results
 - **Asymptotic** normality of MLE
 - **Asymptotic** unbiased of MLE
 - Known sampling variance ...
 - Approximate 95% Confidence intervals:
Find the region of parameter values that yield up to ~ 2 log drop in likelihood ...

Hypothesis testing in a likelihood framework

- Binomial example ...continued
- Q: How to test against the Fisherian $\frac{1}{2}$ sex ratio?
A: compare maximum likelihood with likelihood under the “model” $p=1/2$.



The likelihood ratio test (LRT)

- Idea : comparing the fit of competing (**nested**) model to the data via their likelihood
- LRT statistic
 - 2 competing models M_a and M_b with M_b nested in M_a
 - $G = 2[\text{LogL}(M_a) - \text{LogL}(M_b)]$
- What is the “null” distribution of Λ ?

If M_a is correct $G \sim \chi^2_n$

n is the difference in the number of free parameters fitted in M_a and M_b

Comparing non nested models

- Akaike's Information Criteria (AIC)
- Collection of Models
 - M_1 n_1 free parameters
 - M_2 n_2 free parameters
 - ...
- Compute AIC for each **fitted** model:
 $AIC_i = - 2 \text{ Log}[L(M_i)] + 2 n_i$
- Choose model with **lowest AIC**
- Perspective: robust estimation with **model averaging** ...

Some good reads...

- **Lynch, M. and B. Walsh, *Genetics and Analysis of Quantitative Traits*. 1998, Sunderland: Sinauer Associates, Inc.**

Appendix 4 of this book is a good introduction to likelihood that reviews most of the general results that I find useful when analyzing real data. A bunch of worked simples examples. Connection between LT and te so called G test for goodness of fit

- Burnham, Kenneth P., Anderson, David R. 2nd ed. 2002 **Model Selection and Multi-Model Inference A Practical Information-Theoretic Approach** Springer Verlaag ISBN: 0-387-95364-7

Good introduction to AIC and model selection/averaging

- **Likelihood AWF 1992 Edwards J. Hopkins U Press 2nd edition**

Old fashioned especially in the style and notations but this book was for me the only gateway to the more technical likelihood literature. A must if you enjoy the British academic style !