Maximum Likelihood Estimation Using RTMB

TMB through RTMB

Click here to view presentation online

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Outline: Part I

- Introductions
- Syllabus and our assumptions on background
- Course organization
- Philosophy and approach
- Expectations
- Break

Course intro

- Welcome
- Housekeeping
  - Zoom, github, lecture recordings, communication, etc.
- Round-table introductions
  - Background, why are you here

Syllabus

- Everything through Github
  - https://github.com/QFCatMSU/MLE-Software/
- Syllabus, presentations, code
- If you need help:
  - bence@msu.edu, cahill11@msu.edu

Course organization

- This is a hybrid couse, with some participants in person and others online
- Each class session will be a mix of lecture and work on exercises
- For online participants, exercise work will be in breakout rooms
- No formal homework

Our assumptions about your background

- Some previous experience with statistics including basic idea of fitting models to data
- Some previous use of R
- Some programming (understanding of functions, loops, conditional statements)
- Experience interpreting graphs
- You can get by without all this background, but you should expect to put in more time

Jim's background

- BS in Biology University of Notre Dame
- PhD in Ecology and Masters in Statistics at Univ CA, Santa Barbara (both

1985.

- Worked on evaluation of environmental effects of San Onofre Nuclear Generating Station 1985-1989
- Mathematical Statistician (Stock Assessment Scientist) at NMFS-NOAA Tiburon Lab 1989-1994
- Faculty member at Michigan State University 1994- (retired from tenure stream position July 2023 - currently part time)

Chris's background

- BS. University of Wisconsin Stevens Point (Fisheries science) 2011, MS. University of Alberta (Ecology/Evolutionary biology) 2014. PhD. University of Calgary (Ecology/Evolutionary biology) 2021
- Currently leading efforts to modernize and improve Great Lakes fisheries stock assessments.
- Expertise in hierarchical, state-space, and spatiotemporal modeling and quantitative tools used to inform resource management.
- Associate director of Quantitative Fisheries Center, MSU

Our philosophy on statistical modeling

- Cookbook solutions rarely are adequate for real quantitative problems needed in ecology and resource management
- In class we will solve simple problems the hard way
  - This will make solving hard problems easier and position you to produce better solutions (Royle and Dorazio 2008)
- If you cannot write out your model you don't know what you did!
- Don't get lost in coding.
  - A good model you understand is critical.

Software, implementation, website

- We are primarily going to use R and RTMB
- Recommend Rstudio
- Lectures and code will be available through GitHub
  - You do not need to know how to use GitHub, but that is where you can find code and presentations

GitHub

quarto®

DTU Aqua
National Institute of Aquatic Resources

What is RTMB, why use it?

- R package for maximum likelihood fitting of arbitrarily complex models that incorporate random effects
- Nonlinear and non-normal models (within reason!)
- RTMB uses Template Model Builder (TMB) and TMB was inspired by AD Model Builder (ADMB)

What is RTMB, why use it? A peak under the hood

- Automatic differentiation (as in ADMB)
- Laplace approximation to integrate out random effects (as ADMB)
- Automatic identification of parts of models that are connected (TMB/RTMB only)
- RTMB is much faster than ADMB for models with random effects
- No need for C++ coding (unlike ADMB)
- Major limitation – likelihood must be differentiable function of parameters and RTMB must "see" all the calculations in R

Expectations

- Be kind
- Make an honest effort to learn this stuff
- Share your code
- This course lies at the intersection of mathematics, statistics, ecology, and numerical computing
  - Failure along the way is okay and to be expected and helps you learn and progress

Some disclaimers

1. Maximum Likelihood Estimation is powerful but not without drawbacks and limitations
2. This is not a mathematical statistics course
3. We don't know everything
   - We will do my best to track down answers
4. Please ask questions

Outline: Part II

- Overview of statistical modeling and MLE
- Building blocks
  - Probability, events, and outcomes
  - Random variables and random variates
    - Discrete versus continuous
  - Probability mass and density functions
  - Some common distributions
- The Likelihood function and maximum likelihood estimation

A brief introduction to statistical modeling

- This class is about model-based inference
  - Focus on the development of arbitrarily abstract statistical models
- These models always contain:
  - Deterministic (i.e., systematic) components
  - Random (i.e., stochastic) components
- We estimate model parameters - can be of direct interest, or used to calculate something of interest
  - We are interested in uncertainty of estimates (SEs and CIs)

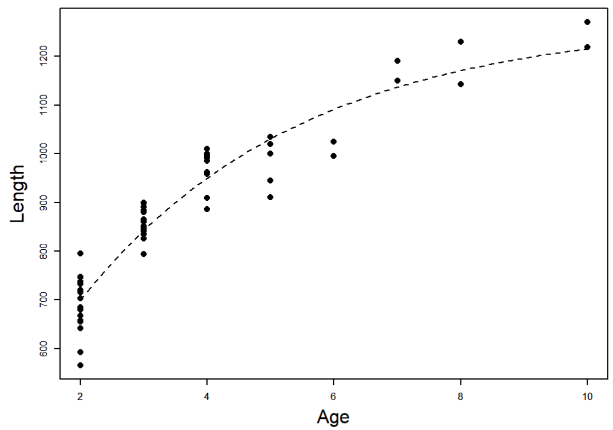
Several example statistical models

- Regression model
- Hierarchical linear model
- Complex age structured assessment model

Regression model example

$$y_i = f(\underline{\theta}, \underline{X}) + \varepsilon_i$$

$$L_i = L_\infty \left(1 - e^{-K(a_i - t_0)}\right) + \varepsilon_i$$



Data: T. Brenden unpublished. Photo: E. Engbretson, USFWS https://commons.wikimedia.org/w/index.php?curid=3720748

Hierarchical linear model example

- Weight is power function of length multiplied by error
  - On log scale the relationship is linear with additive error
- i represents ponds, j fish within ponds
- Intercept and slope vary randomly among ponds, residual variance is pond specific
- Y~N(a,b) means Y is normal, with mean a and variance b

State Space Catch at age model

State Space Catch at age model (continued)

Probability

- Whole books about definitions and meaning.
- I follow a frequentist definition for intuition, while recognizing that there is some logic to Bayesian claims of degree of belief
    - Frequentist definition: The long run proportion of of times an event occurs under identical conditions
    - Statisticians sometimes distinguish outcomes from events. Outcomes are really elementary events. Event might be catching a fish in 7 to 8 inch bin, outcome would be catch a fish and measure its length.

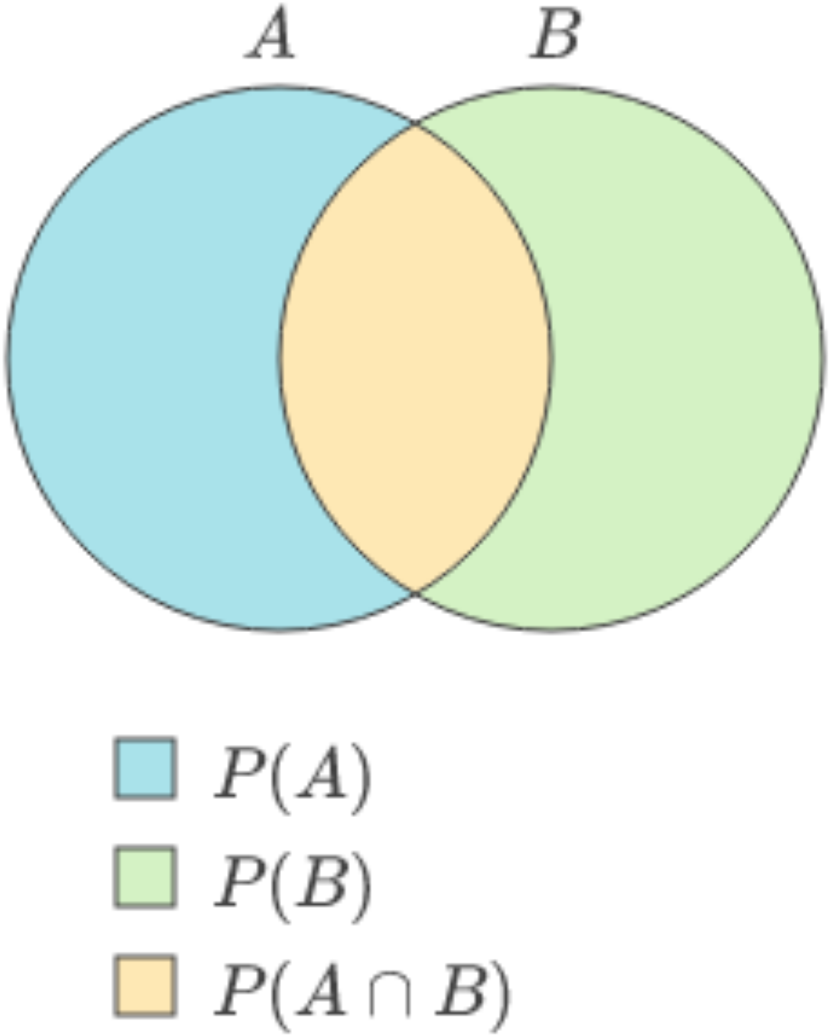
Basic properties (axioms) of probabilities

- The sum of probabilites over all possible mutually exclusive events is 1.0 (So something will happen)
- Probability of any given event is 0 (and 1)
- The probability of the union of mutually exclusive events is the sum of their separate probabilities
  -
- if A and B independent

Conditional probability

- "|" read as "given or conditional on – Probability of A given B
- Conditional probabilities recognize that the occurrence of event B can provide information on whether event A will occur
- Convince yourself that if A and B independent

Conditional probability



Conditional Probability Formula

$$P(A \mid B) = \frac{P(A \cap B)}{P(B)}$$

Probability that A occurs given that B has already occured

stats.stackexchange.com/questions/587109

Random variables (words)

- Technical definition is that they are functions that convert probability spaces for events/outcomes to numeric results.
  - Ironically they are neither random nor variables!
- Less technically (but still techno speak!) they describe the numeric outcome of a random process. I.e., they are not a number (or vector/matrix of numbers) but rather the process of producing them.
- A random variate is a particular numeric outcome
- Text books say usually capital letters used for random variables and lower case for random variates.

Random variables: math expression for simple example (coin flip)

We flip a coin and call a heads 1 and a tails 0:

- Pr = probability
- We could say the random variable Y has a Bernoulli distribution

Bernoulli probability mass function (pmf)

- The pmf is a function that calculates the probability given the random variate (value) and the parameter(s) (here)

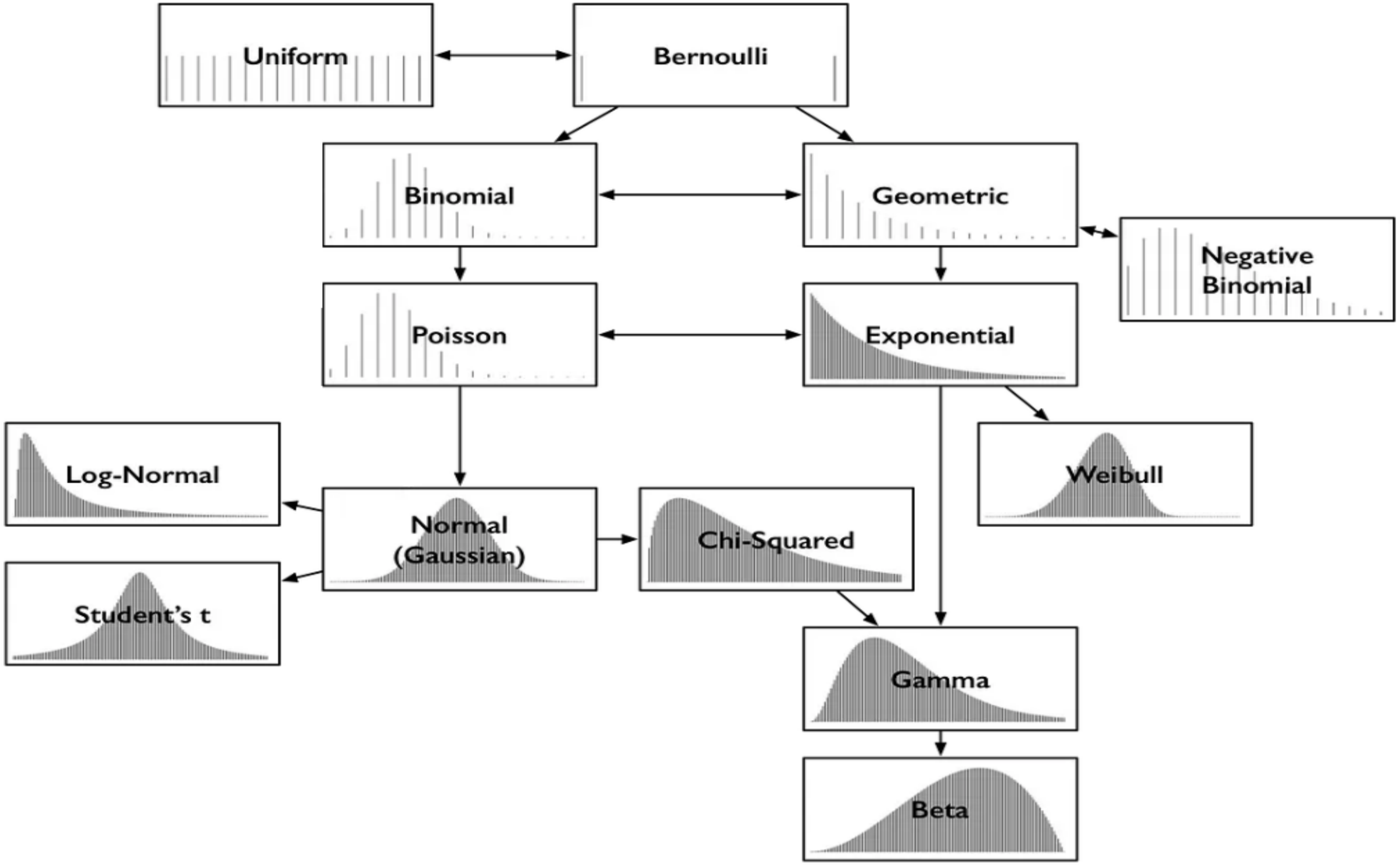
  is observed datum

  - pmf for discrete outcomes
- If this was a continuously distributed random variable we would use probability *density* function (pdf)

pmf and pdf notation

- A conventional notation for this stuff is
- Sometimes with subscript for random variable:
- Conditional bit indicates that the probability of an observed value depends on parameter(s) used to specify the distribution of the random variable
- Notation for Benouilli random variable:

Some common statistical distributions



Common probability distributions and some key relationships

Figure Reference

More notation notes for everyone's sanity

- In general we will provide the pmf (or pdf) expressed as a function of and the parameters of the distribution.
- For example, will use to indicate a random variable is normally distributed with mean and variance
- In general regular font for scalars, bold for vectors and matrices

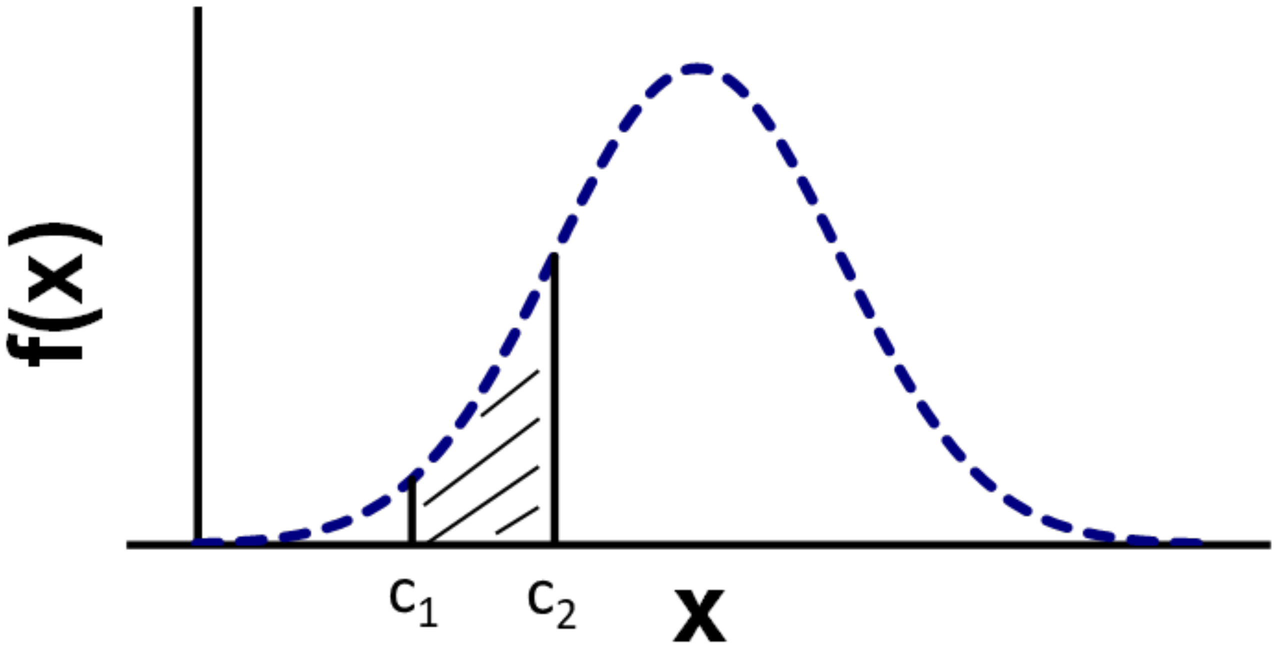
More notation

- You might see it this way too:

Discrete versus Continuous random variables

- Discrete means the set of possible outcomes is countable with each possible value having an associated probability (calculable from the pmf).
- Continuous means not countable (generally this means there are infinite numbers of possible values between any two other possible values). Pr(y) for any particular y is 0. So we use a probability density function.
- Intuition/common sense sometimes used to choose between the two. E.g., catch or CPUE often modeled as continuous

Probability density function



- Pr()=Pr()=0 !
- Area under the pdf function gives probability for interval
- Pr()=Pr()-Pr()

Cumulative distribution function

- $F(x)=Pr(X<x)$
- For continuous variables, the derivative of $F(x)$ with respect to x is $f(x)$ (the density)
  - Why? Does this make sense?

Joint probability density and mass functions

Vector of observed values, with elements having the same pdf/pmf or different ones and these random variables might be independent or not:

Special case of each element representing an independent random variable:

Special special case of independent and identically distributed (iid) random variables:

These special cases very important for practical MLE work!

The likelihood function

- No new math!!!
- The likelihood function is just the joint pdf re-expressed as a function of the parameters:

Maximum likelihood estimation

- Adjust until is maximized
- The rest is “just” details :->

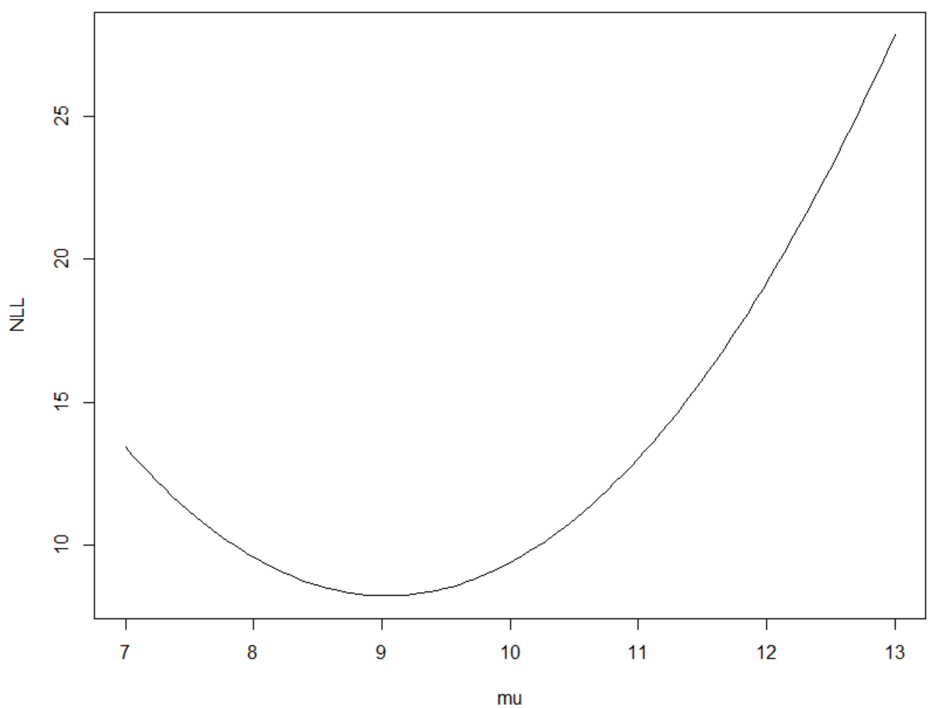
A numerical example of joint likelihood

- **x** ={10.72,7.23,10.07,8.62,8.55}
- Each observation (x) independent from a common normal distribution with mean 10, variance 2 (i.e., they are iid)
- Calculate the likelihood of these data, i.e., f(10.72)f(7.23)f(10.07)f(8.62)f(8.55) using R
- Hints. The result is just a single number. You can calculate the pdf of f(x) for a normal distribution in R using the dnorm function. The dnorm function uses sigma (SD) not sigma squared (variance)
- Time permitting - generalize your solution as a R function to calculate likelihood for any vector x, with specified mean, variance.

Working with the log likelihood (prefered for numerical reasons)

- Perhaps obviously, if you adjust parameters to maximize the log of the likelihood function this will also maximize the likelihood.
- RTMB and most software minimizes the negative log likelihood rather than maximizing log likelihood (convention)
- Working on the log-scale improves numerical performance.
    - The joint log likelihood for independent observations is the SUM (rather than product) of the individual log f(x) values

Negative log likelihood versus mu for five iid observations from a Normal distribution (known variance of 2)



Detection probability example - Sneak turtles

- 30 turtles released in pond two recaptured later
- Release is trial, recapture is a success
- If each trial is independent, number of successes binomial
- Just one observation - y successes. Likelihood just the pdf

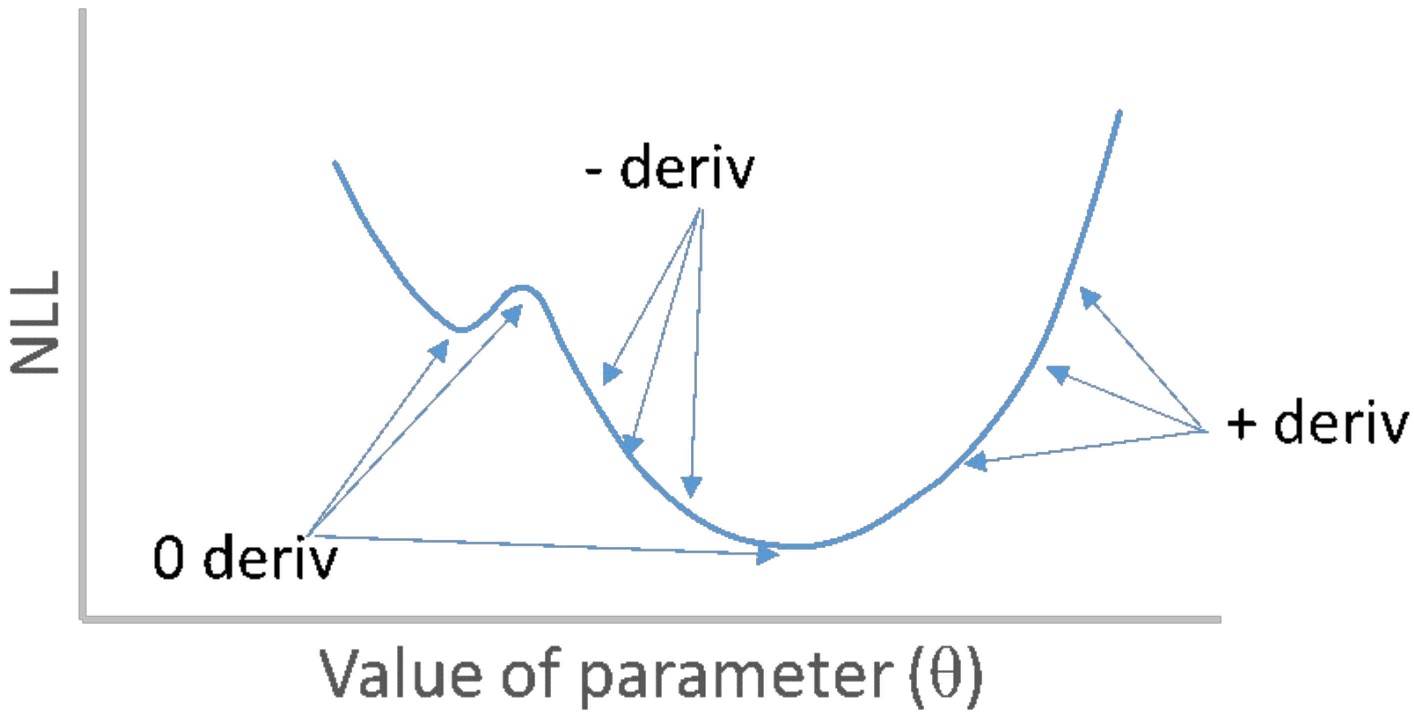
Finding the MLE - The sneaky example

- Determine the detection probability of invasive Alabama sneak turtles by releasing some into a small pond and counting them on a later occasion
- We release n = 30 and later count y = 2 sneakers
  - successes vs. trials
- What's probability of detecting a turtle, i.e., ?

Methods for finding MLEs

- Analytical solution (involves derivatives)
- Grid search
- Iterative searches
  - Non-derivative methods
  - Derivative methods (such as quasi-Newton)

Role of derivatives in finding MLEs



NLL as function of single parameter with derivatives

- Derivatives of NLL with respect to parameters zero at minimum
- Second derivatives of NLL with respect to parameters are positive at minimum

Sneak turtle analytical approach

Grid search for sneak turtle example

```
y <- 2 # successes
n <- 30 # trials
eps <- 1e-6 # buffer because theta must lie within [0,1]
theta <- seq(eps, 1 - eps, length.out = 1e4) # sequence to thetas to try
 # plug thetas into pmf
logLike=y * log(theta) + (n - y) * log(1 - theta)
MLE <- theta[which.max(logLike)] # maximum likelihood estimate
my_data <- data.frame(logLike, theta) # for plotting if you want
```

Analytical vs. grid search MLEs

```
y / n # analytical theta MLE
```

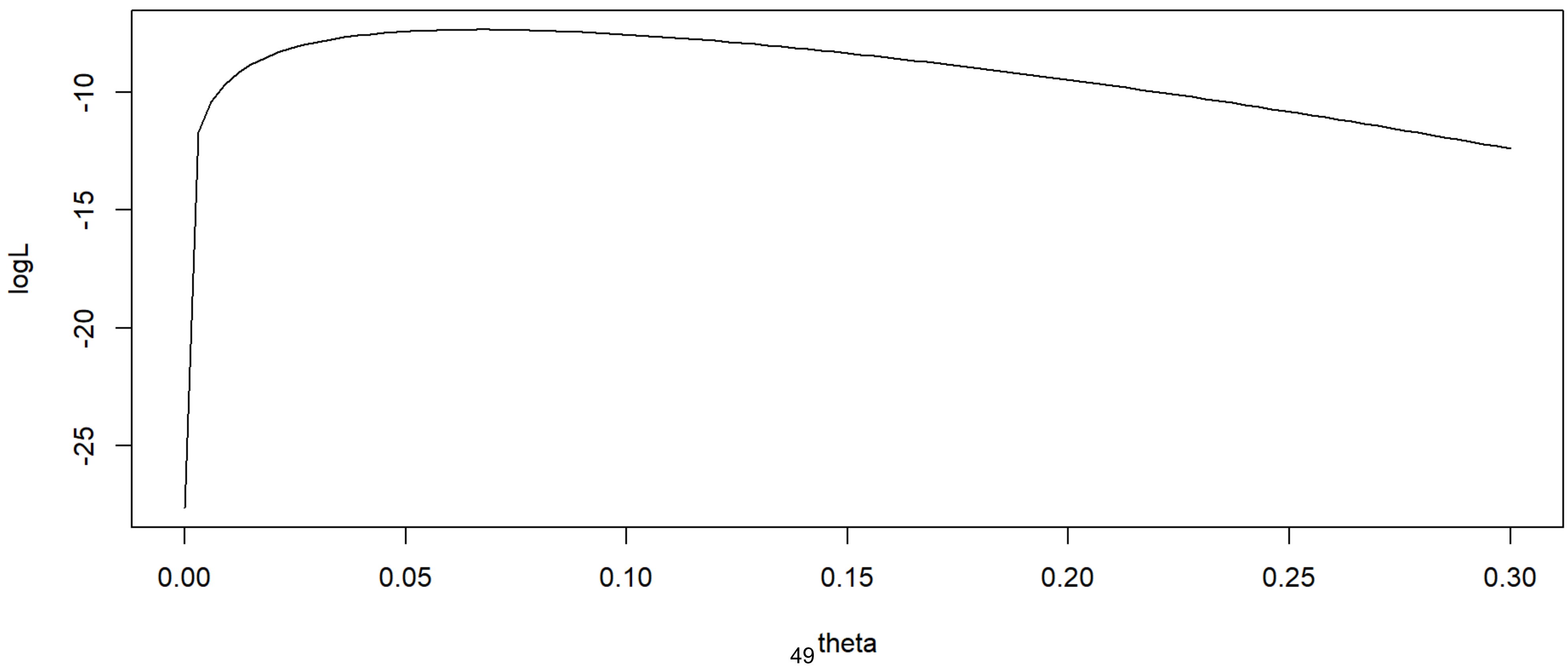
```
[1] 0.06666667
```

```
MLE # grid search approximation
```

```
[1] 0.06670754
```

Plot of log likelihood function for sneak turtle

```
curve(y * log(x) + (n - y) * log(1 - x),from =eps, to=0.3,xlab="theta",ylab
```



Exercise - use built in binomial distribution

- Calculate the log likelihood at the MLE using logLike = dbinom(y,n,MLE,log=TRUE)
- Compare with logLike = y * log(MLE) + (n - y) * log(1 - MLE)
    - Why are they different?
- Steal the grid seach code, run it.
    - Then replace logLike = y * log(theta) + (n - y) * log(1 - theta) by logLike=dbinom(y,n,theta,log=TRUE)
    - You should get the same grid search estimate

Another exercise or interest of time demo

- For the previous sample of five observations from normal find the MLE estimate of the mean assuming the variance known equal to 2 by conducting a grid search
- Think about how you would do grid search to find MLE estimates of both the mean and variance at the same time

The regression case

- Observations assumed independent but not identically distributed.
- The mean varies among observations and in this simple case variance is same for all observations:
- Cannot estimate mean as a parameter for every observation
- But can calculate it as function of parameters, e.g, :
- general message estimated parameters are not the same as the distributional parameters for the pdfs/pmfs
- What are the estimated (model) parameters?

Psuedocode for the regression problem

- Specify , , and
- Calculate
- Calculate NLL

Search over different values of , , and and repeat 1-3 until you find the values that minimize the NLL

Two ways to frame the regression model

- Previously we modeled mean, which was a distributional parameter. Now we write a model for the individual observations. We can write the likelihood in terms of the errors, or the observations - The two are theoretically equivalent!
- Note but don't worrying about. The standard model notation uses random quantities on the RHS but specifies the random variate on the LHS

Properties of MLEs

- Terminology: ML Estimator versus ML Estimate
- Ideal estimator is lowest variance among unbiased estimators
- MLE not guaranteed to do this!
- MLEs are consistent, meaning estimates will become closer to correct values and sample sizes increase
  - Asymptotically unbiased
  - Errors get smaller with more data

Familiar example of bias for an MLE

- MLE for variance of normal random sample:
- Expected value:
- Standard (unbiased) estimator: