

IPS-164 INTRODUCTION TO PHYLOGENETICS 2022

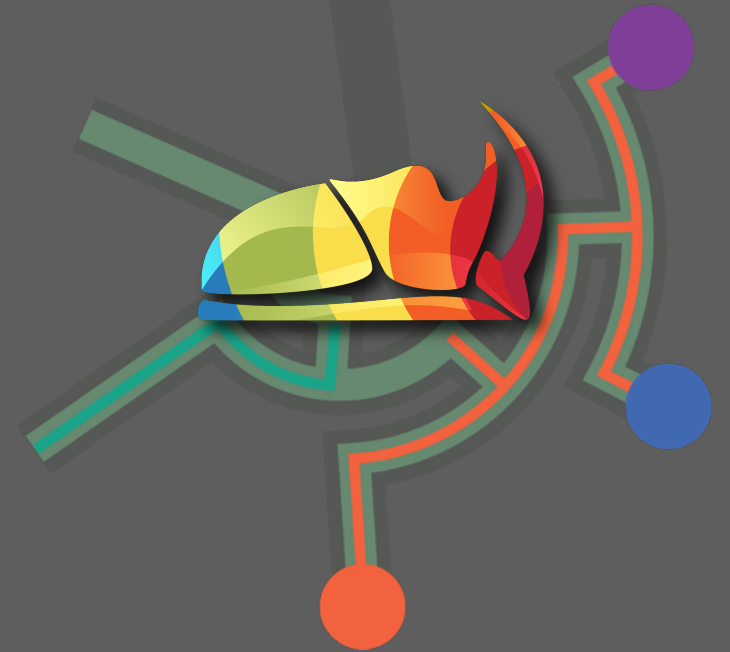
## Lecture 6

# Intro to statistical phylogenetics. Part I

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Finnish Museum of Natural History, University of Helsinki



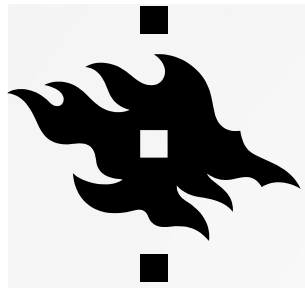
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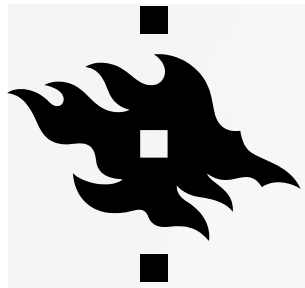
• <https://www.tarasovlab.com>



# LET'S GET TO KNOW EACH OTHER

1. Say your name
2. Your interests in biology/phylogenetics
3. What has brought you to this course?
4. Your expectations from the course?





# PLAN OF THE TODAY'S LECTURE

1. Intro to this (the second) part of the course
2. Overview of the statistical phylogenetics: which questions statistical phylogenetics can address?
3. Parsimony vs. statistical phylogenetics
4. Intro to statistics and modeling
5. Binomial model

# Aim of this part of the course

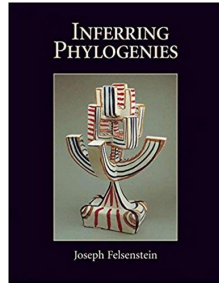
- Explain how the statistical inference works in phylogenetics and overview its main field
- So, you will be able to calculate likelihood by “hand”
- You will learn how to reconstruct phylogenetic tree and perform various other analyses
- You will be able to select amongst available methods to address your research questions



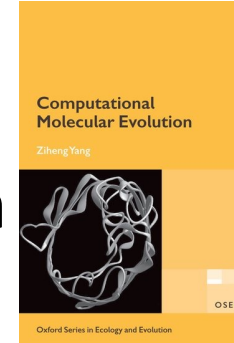
# Lectures in this part of the course

6. Introduction to statistical phylogenetics (part I)
7. Introduction to statistical phylogenetics (part II)
8. Reconstructing phylogenies (part I)
9. Reconstructing phylogenies (part II)
10. Tree dating
11. Trait evolution
12. Trait evolution and Diversification

# Suggested literature

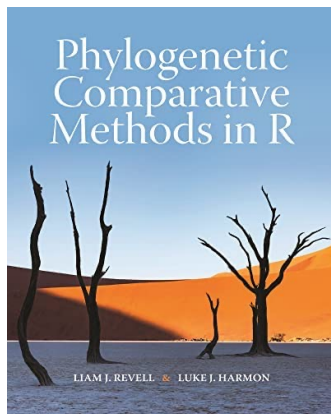
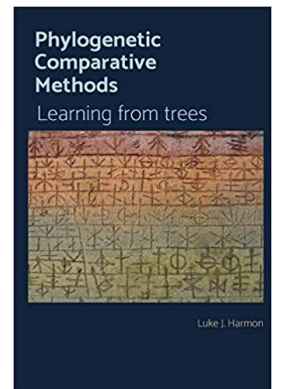


- Computational Molecular evolution
- Felsenstein's Inferring phylogenies



- Luke Harmon. Phylogenetic Comparative Methods

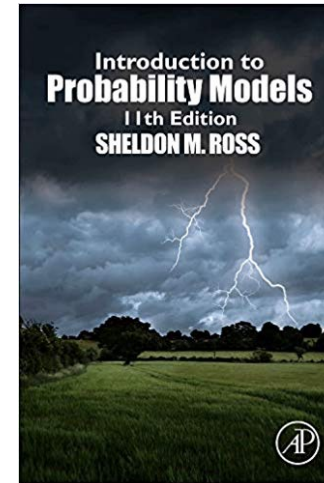
It's free at <https://lukejharmon.github.io/pcm/>



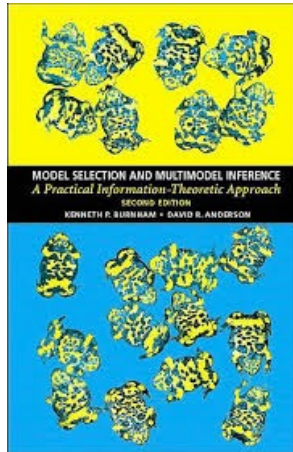
- Phylogenetic Comparative Methods in R

# Suggested literature

- Intro to Probability models



- Model Selection by Burham and Anderson



# Developing skills in Statistical Programming with R or Python

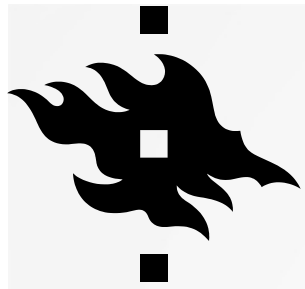
- Books

- Paradis. Analysis of Phylogenetics and Evolution with R
- Revell & Harmon. Phylogenetic Comparative Methods in R
- Grolemond. R for Data Science. <https://r4ds.had.co.nz/index.html>

- All materials will be available at:

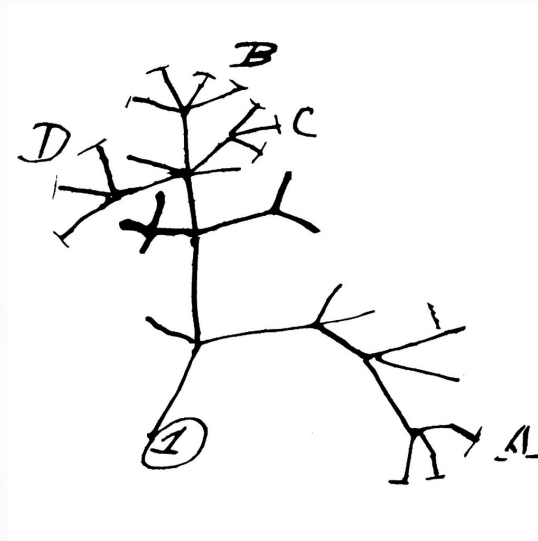
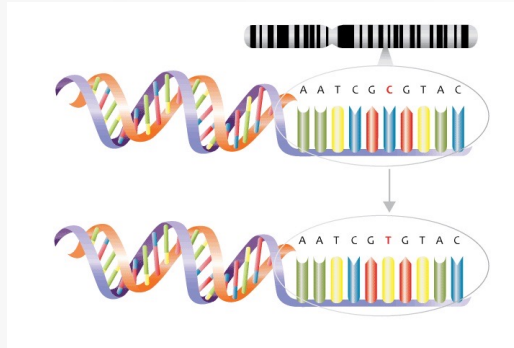
- GitHub [https://github.com/sergeitarasov/Course\\_IPS-164](https://github.com/sergeitarasov/Course_IPS-164)
- UH website <https://www.mv.helsinki.fi/home/jhyvonen/IPS-164/>

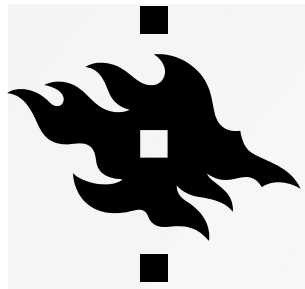




# WHICH QUESTIONS STATISTICAL PHYLOGENETICS CAN ADDRESS

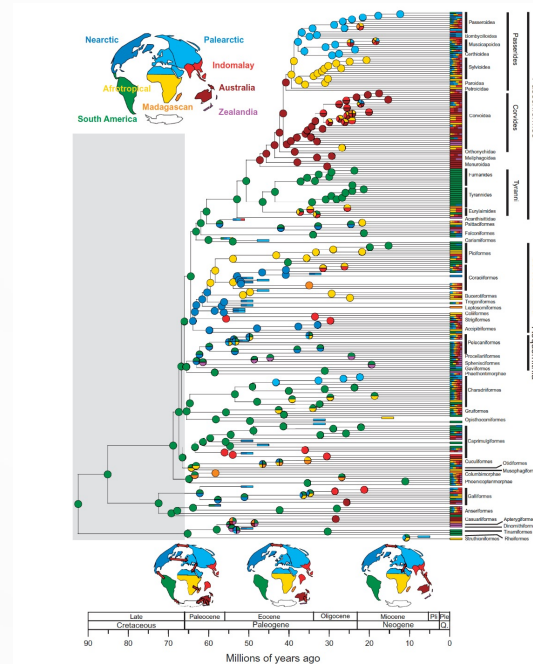
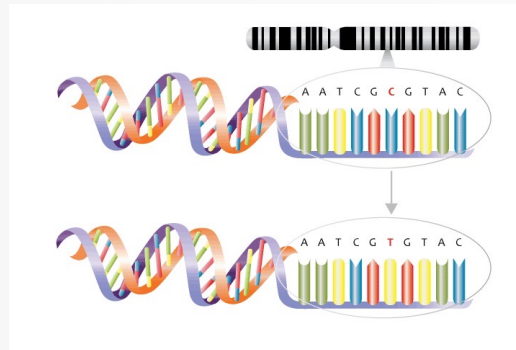
## 1. Tree Reconstruction



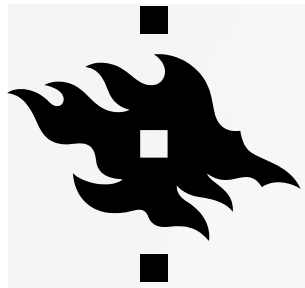


# WHICH QUESTIONS STATISTICAL PHYLOGENETICS CAN ADDRESS

## 2. Tree Dating

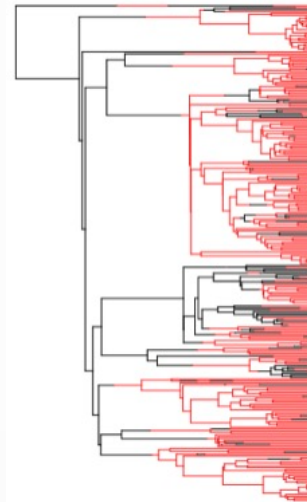
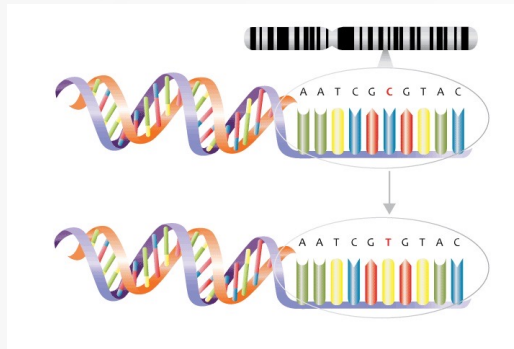


Fossil record



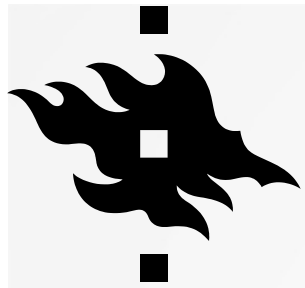
# WHICH QUESTIONS STATISTICAL PHYLOGENETICS CAN ADDRESS

## 3. Ancestral Character State Estimation



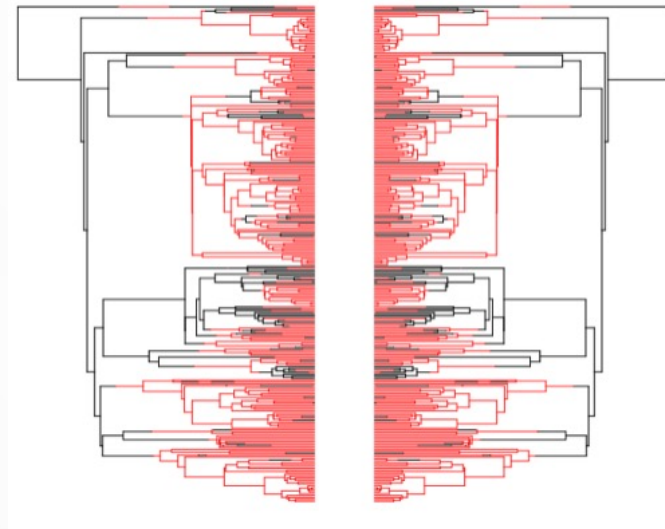
Discrete and  
Continuous traits

Give you example of traits!

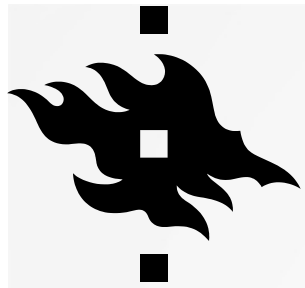


# WHICH QUESTIONS STATISTICAL PHYLOGENETICS CAN ADDRESS

## 4. Correlation Between two or more traits

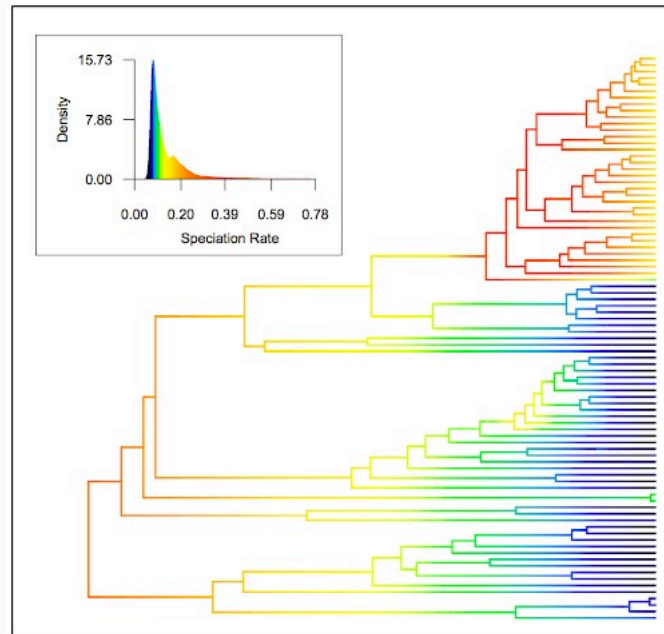


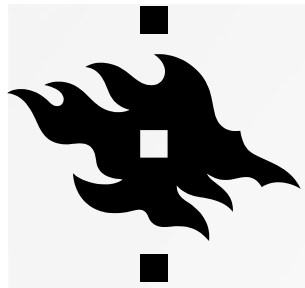
Discrete and  
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# WHICH QUESTIONS STATISTICAL PHYLOGENETICS CAN ADDRESS

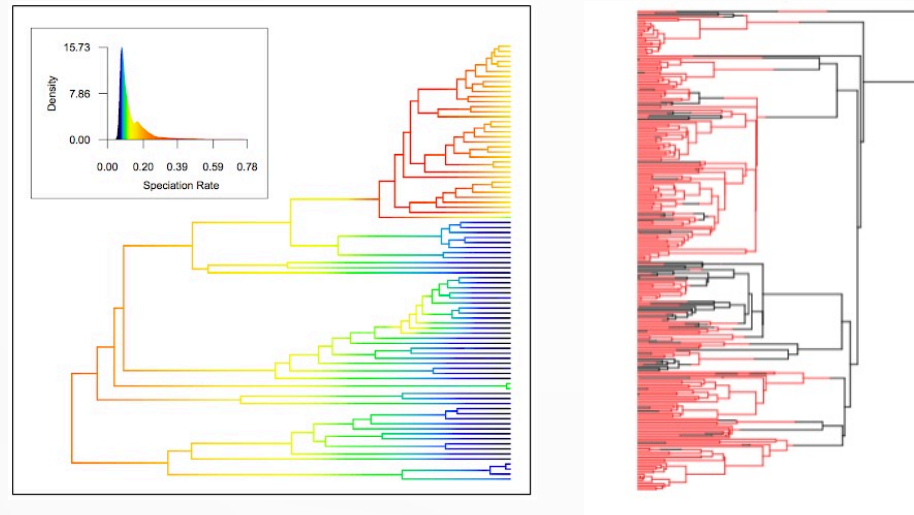
## 5. Reconstructing Diversification process

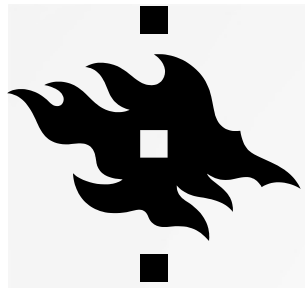




# WHICH QUESTIONS STATISTICAL PHYLOGENETICS CAN ADDRESS

## 6. Correlation between Diversification and traits





# MAIN STATISTICAL METHODS

## Likelihood and Bayesian inferences

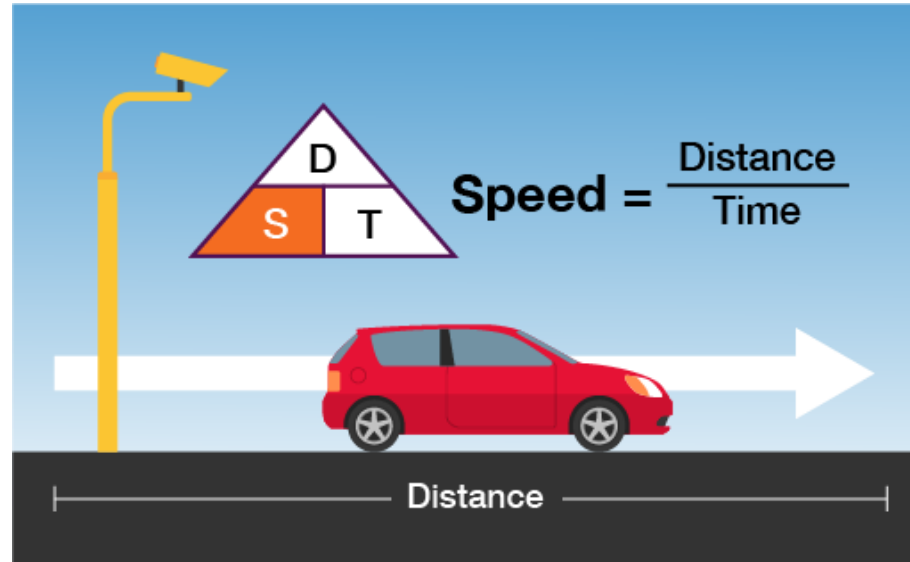


Ronald Fisher



Thomas Bayes

# Modeling natural and phylogenetic phenomena

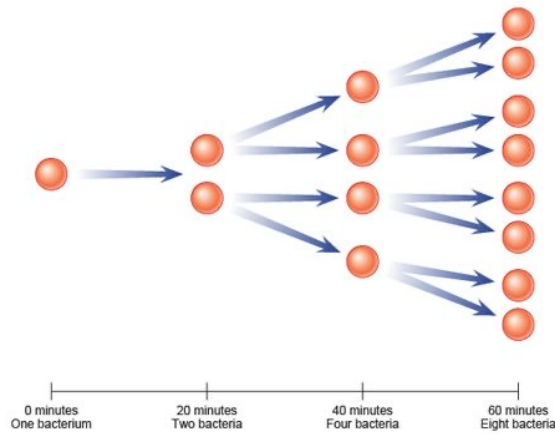


Models are common in physics

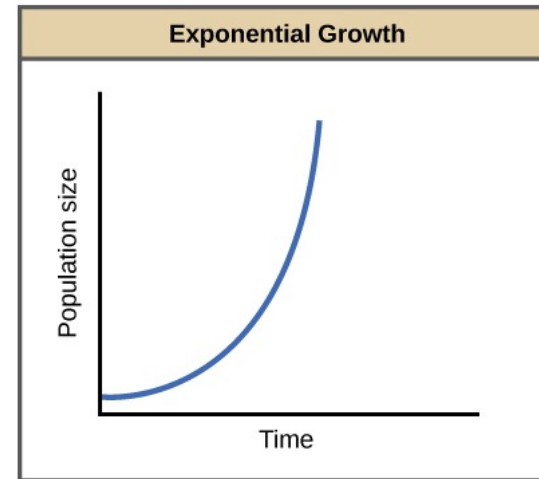
Models are sets of rules describing how a system changes over time



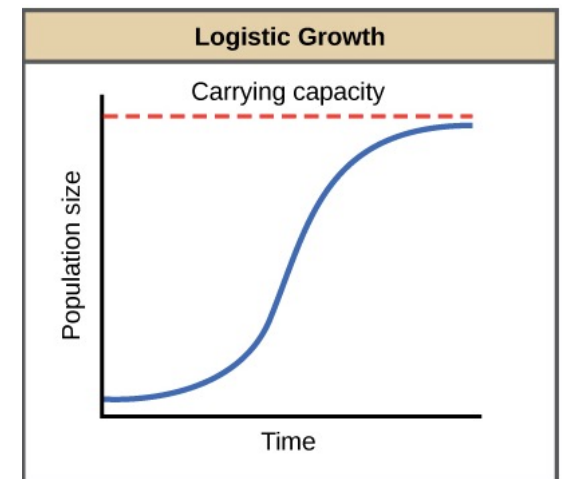
# Modeling natural and phylogenetic phenomena: Bacterial Growth



$$\text{Population size} = 2^t$$



$$\text{Population size} = 2^t$$

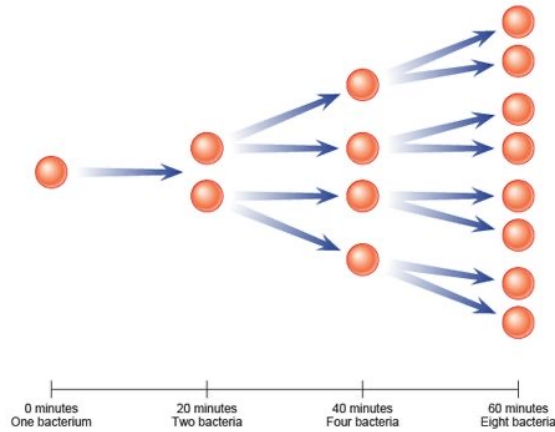


$$f(x) = \frac{L}{1 + e^{-k(x-x_0)}}$$

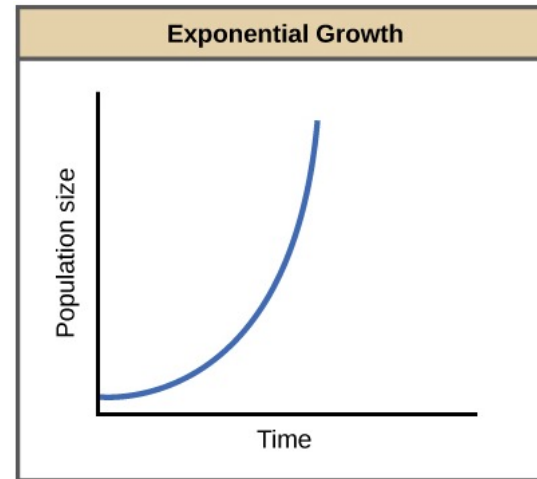
Models are sets of rules describing how a system changes over time

# Modeling natural and phylogenetic phenomena: Bacterial Grow

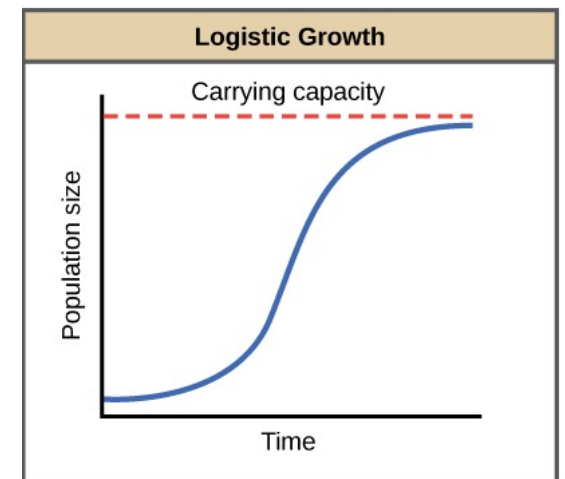
Q: Do you know any other models used in biology?



$$\text{Population size} = 2^t$$

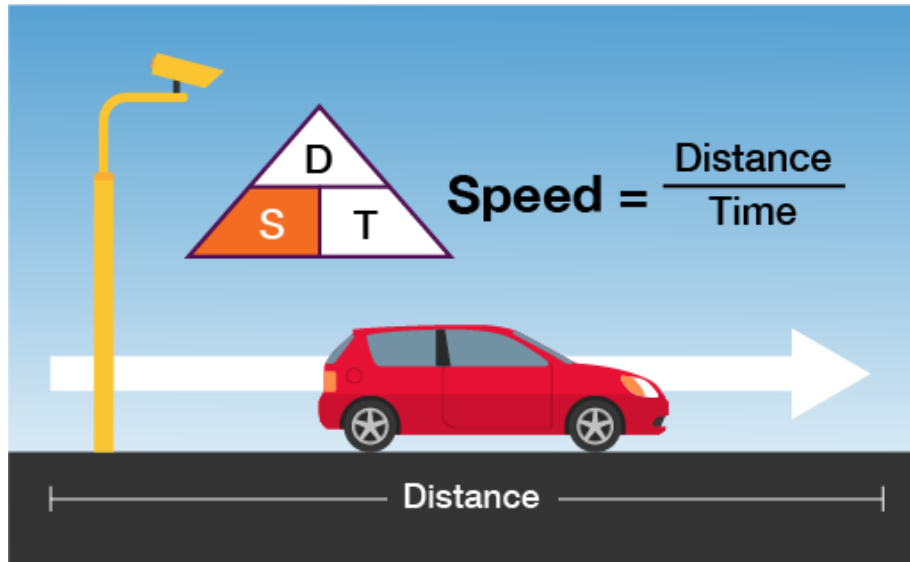


$$\text{Population size} = 2^t$$

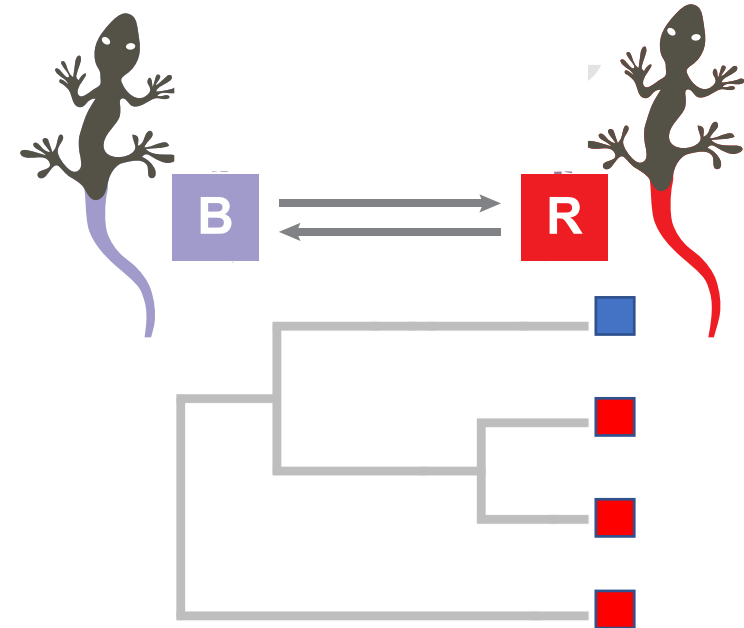


$$f(x) = \frac{L}{1 + e^{-k(x-x_0)}}$$

# Modeling natural and phylogenetic phenomena



Models are common in physics



Models in biology?

Models are sets of rules describing how a system changes over time

# Parsimony vs. Statistical Phylogenetics

**Parsimony**

**Statistical models**

# Parsimony vs. Statistical Phylogenetics

## **Parsimony**

- Strict principle of Occam's razor
- Not a statistical model
- But can be considered as a mathematical model
- Some analysis are challenging (e.g., tree dating)
- FAST

## **Statistical models**

# Parsimony vs. Statistical Phylogenetics

## **Parsimony**

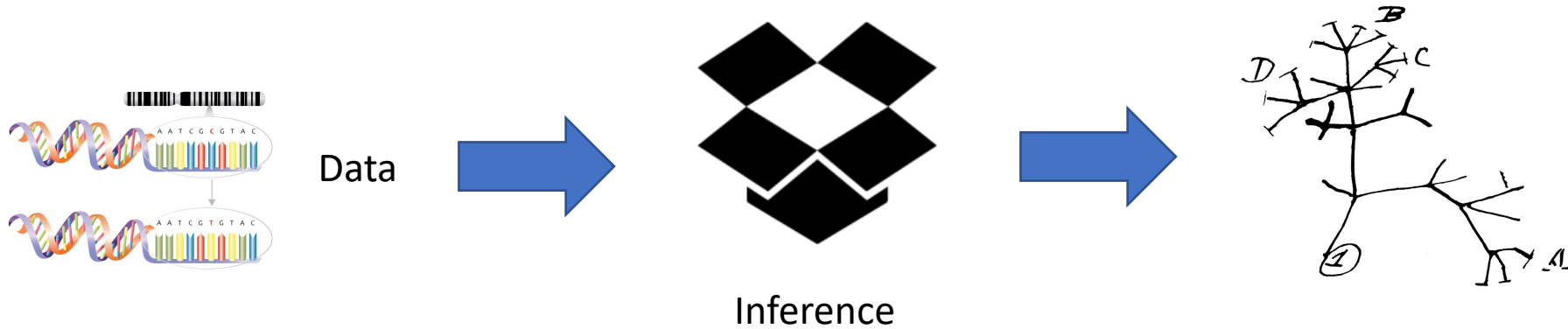
- Strict principle of Occam's razor
- Not a statistical model
- But can be considered as a mathematical model
- Some analysis are challenging (e.g., tree dating)
- FAST

## **Statistical models**

- Can model manifold of natural phenomena & processes
- Many different methods available
- Many different models to create
- Can be SLOW

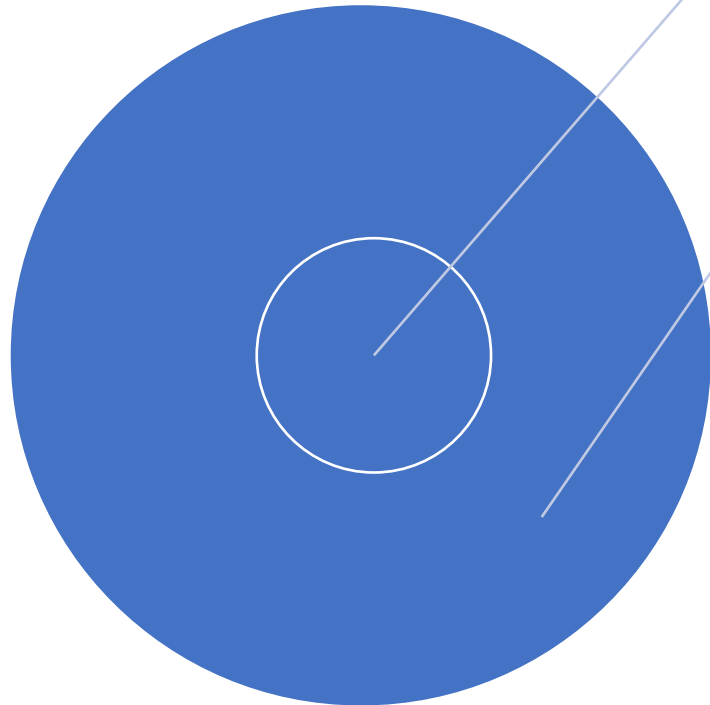
# Aim of this part of the course

- Main principles of modeling data

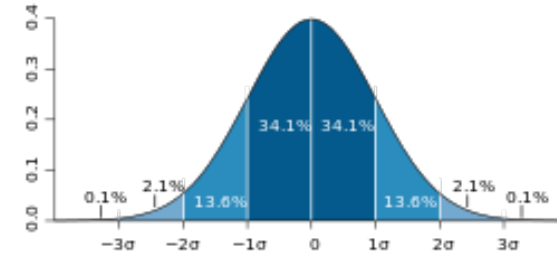


# Modeling principles in phylogenetics

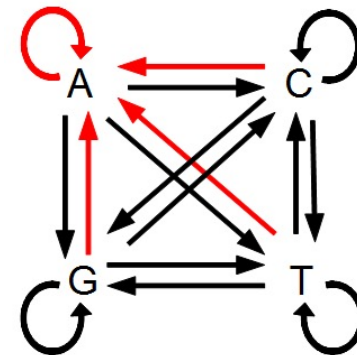
PHYLOGENETICS



Probability Distribution



Stochastic models of character evolution (DNA or Morphology)





# Intro to maximum likelihood (ML) method using simple Binomial model (coin toss)

- Rules of Probability
- Probability Distribution
- Binomial Model
- Likelihood of Binomial Model



# Quick intro to probability: main rules

- AND rule

$$\text{🪙 AND 🪙} = 0.5 * 0.5 = 0.25$$

- OR rule

$$\text{🪙 OR 🪙} = 0.5 + 0.5 = 1$$



- Sum of all events

$$\text{🪙 🪙 🪙} = 1$$

3 trials

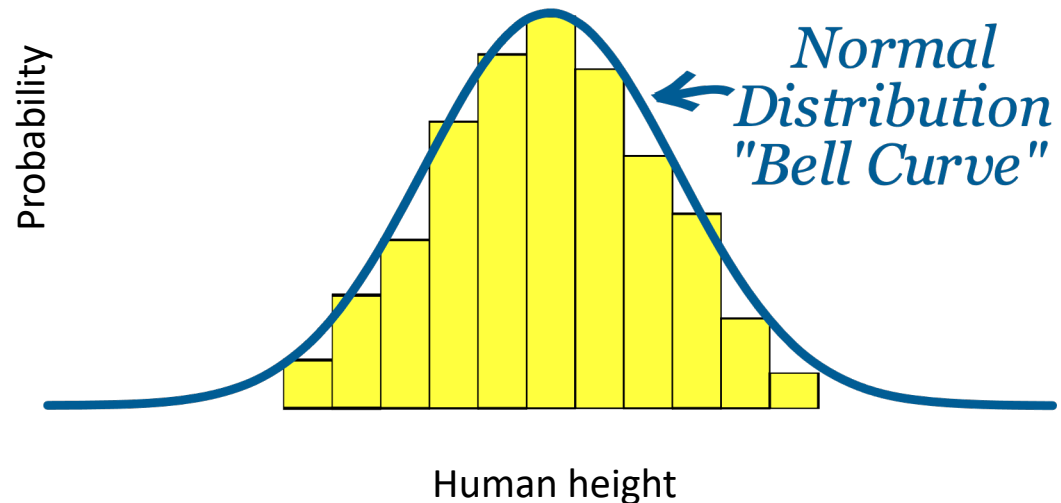
# Let's play with a fair dice.

- Q1: Probability of seeing "1" after one trial?
  - $1/6$
- Q2: Probability of seeing "2" AND "4" given two trials?
  - $1/6 * 1/6 = 1/36$
- Q3: Probability of NOT seeing "3" given one trial?
  - $1 - 1/6 = 5/6$
- Q2: Probability of seeing "1" OR "6" given one trial?
  - $1/6 + 1/6 = 1/3$



# Probability distribution

- A probability distribution is a function that provides the probabilities of occurrence of different possible outcomes in an experiment.
- Distribution usually refers to a distribution of a random variable
- In probability and statistics, a random variable, is a variable whose possible values are outcomes of a random phenomenon.

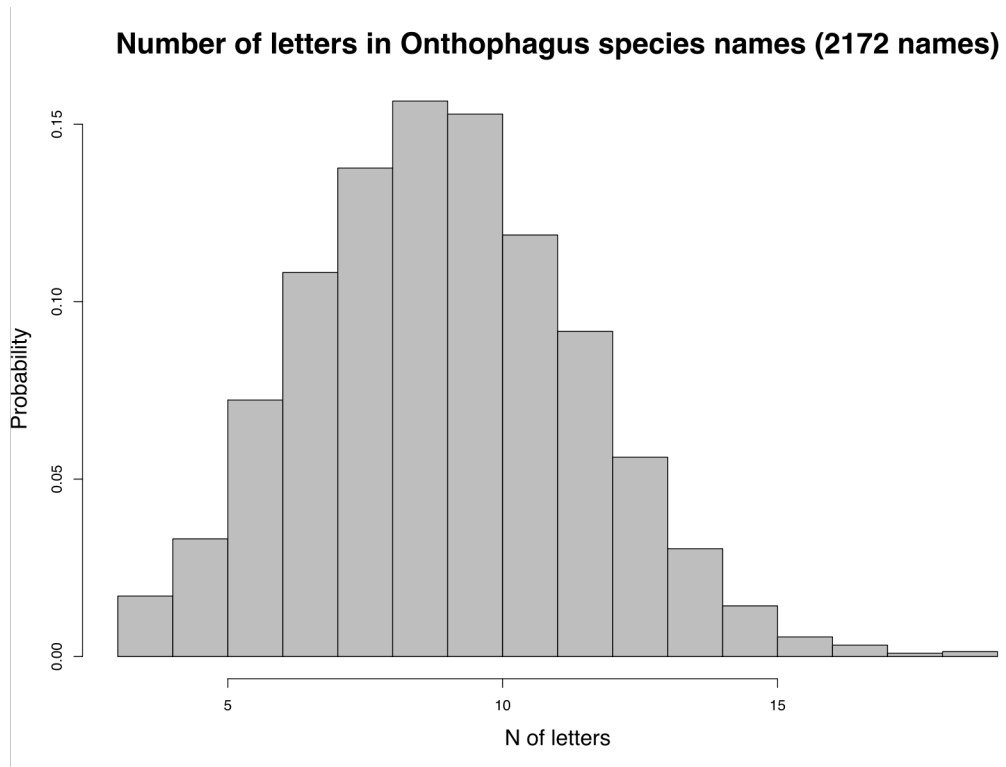


# Informal Axioms (rules) of Statistics

- Any measured quantity of any set of objects in the Universe has some probability distribution
- There are ~20 most common distributions in the Universe (e.g., Binomial, Normal, Gamma, Poisson etc.)
- Most likely, the measured quantity falls into one of those ~20 common distributions

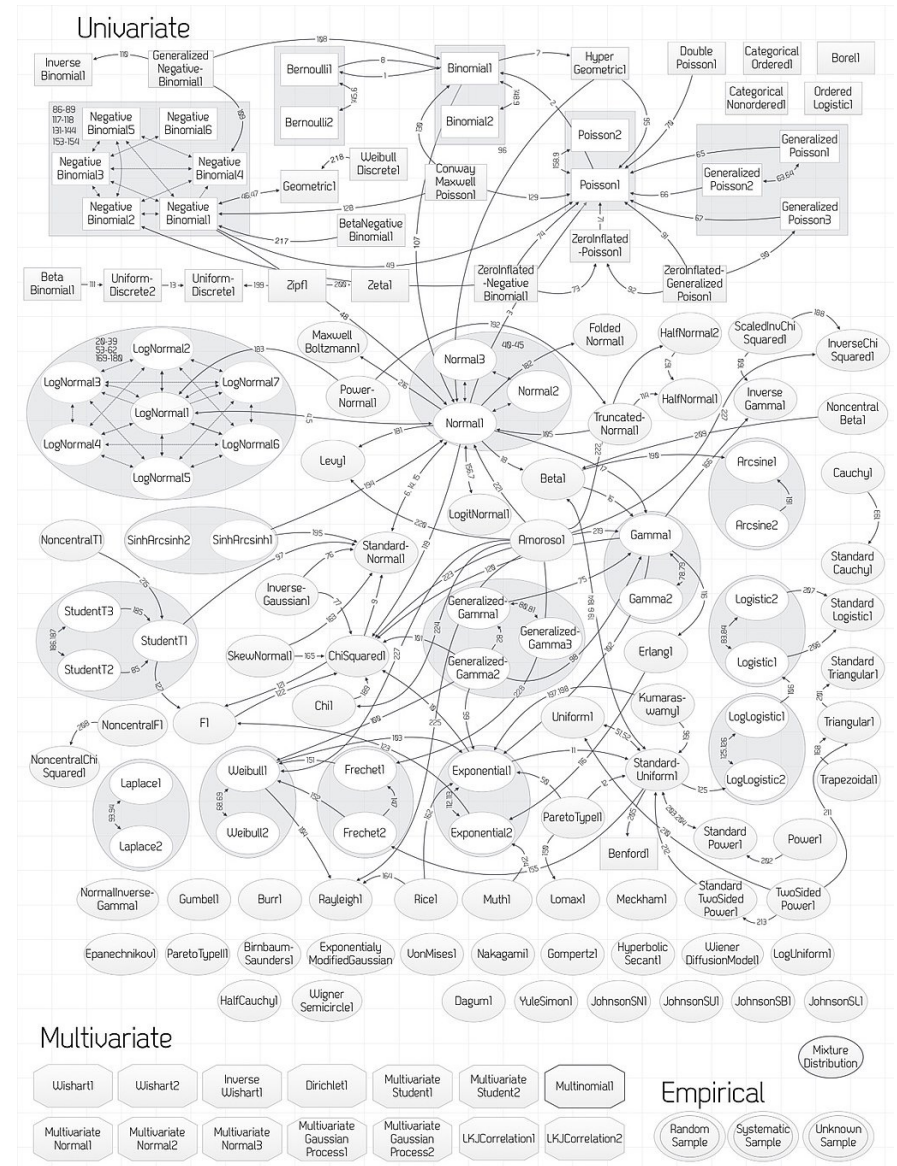
# Empirical probability distributions

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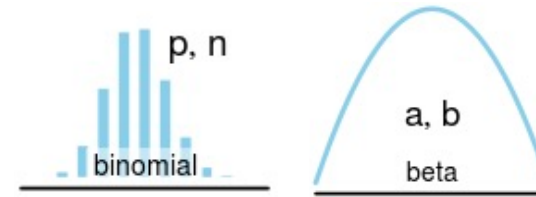
# Relationships among probability distributions

[https://en.wikipedia.org/wiki/Relationships\\_among\\_probability\\_distributions](https://en.wikipedia.org/wiki/Relationships_among_probability_distributions)

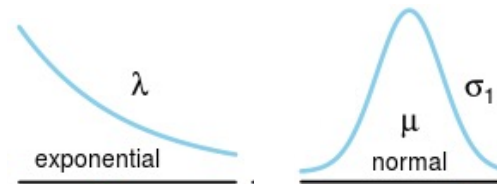


# Classification of probability distributions

- Discrete vs. Continuous

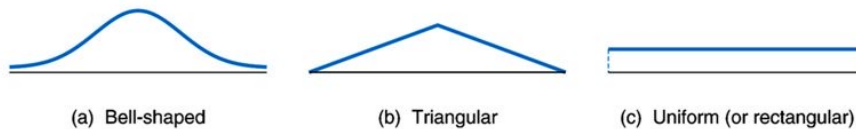


- By number of parameters

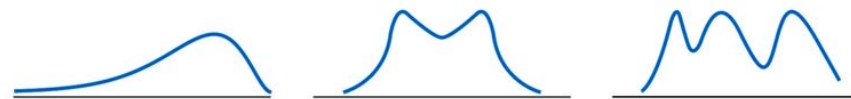


- By domain  $[-\infty, +\infty)$  vs.  $[0, +\infty)$  vs.  $[0,1]$

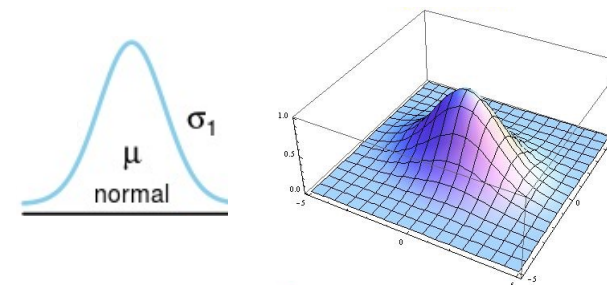
- By shape



- By mode: unimodal vs. multimodal



- By dimension of random variable: univariate vs. multivariate





# Binomial model ( and distribution)

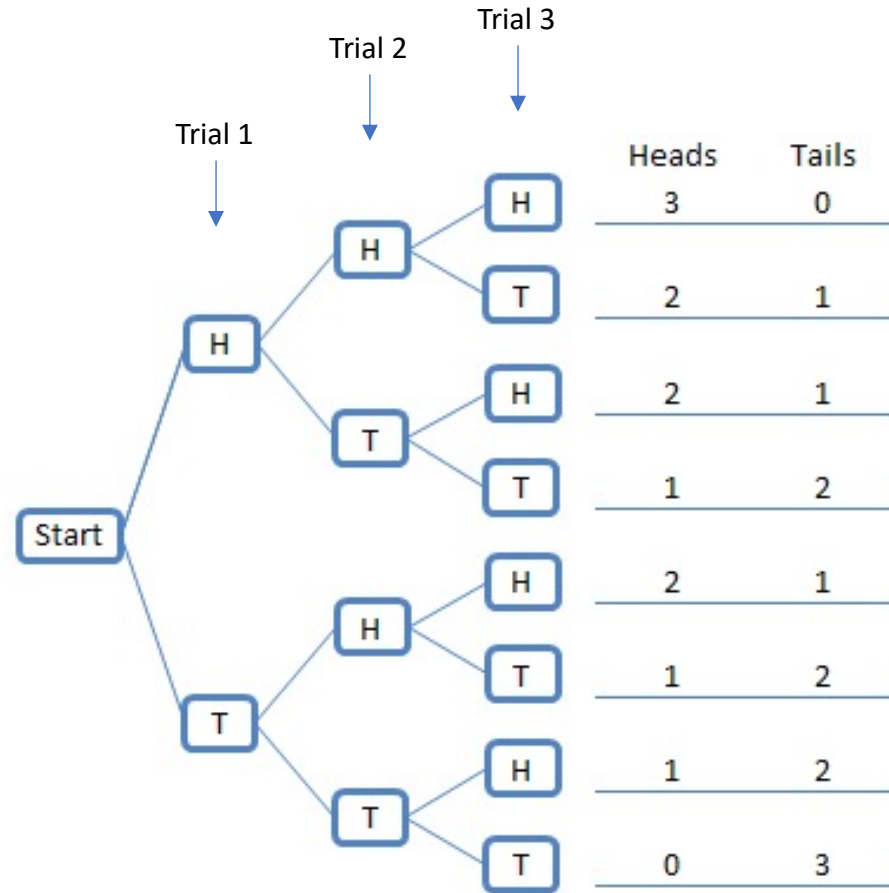
Binomial model gives the probability of seeing  $k$  heads in  $n$  coin tosses (trials) given that probability of seeing a head in one coin toss is  $p$ .

Let's consider an example where  $n=3$  and  $p=0.5$

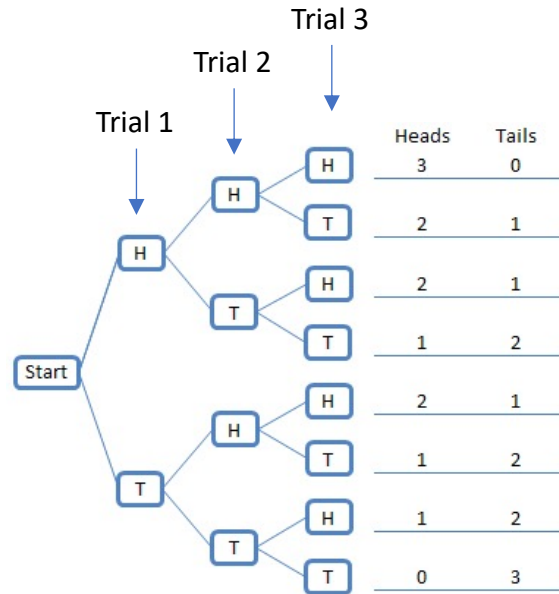



- Coin is fair
- We toss the coin 3 times


# Estimating number of heads



# Estimating probabilities



 =  $0.5^3 = 0.125$

 =  $0.5^3 = 0.125$

..... =  $0.5^3 = 0.125$

# Estimating probabilities

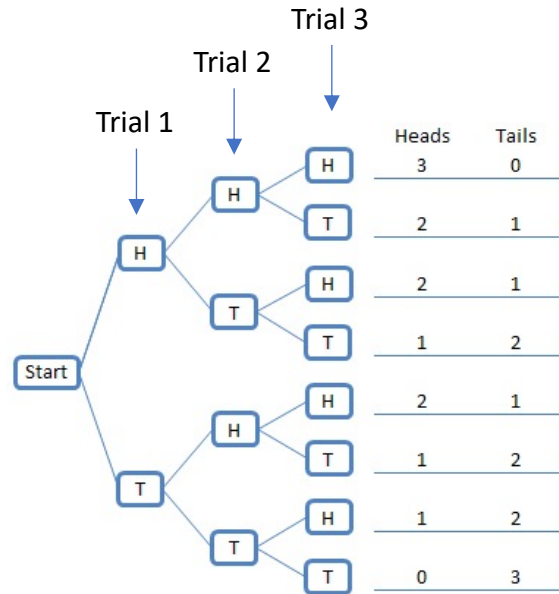
$$\binom{n}{k} = \frac{n!}{k!(n-k)!}$$

$$\binom{n=3}{k=3} = \frac{3!}{3!(3-3)!} = 1$$

$$\binom{n=3}{k=2} = \frac{3!}{2!(3-2)!} = 3$$

$$\binom{n=3}{k=1} = 3$$

$$\binom{n=3}{k=0} = 1$$



$$= 0.5^3 = 0.125$$

$$= 0.5^3 = 0.125$$

.....

$$= 0.5^3 = 0.125$$

# Estimating probabilities

Number  $k$  heads in  
3 trials

Probability of seeing  $k$  heads  
in 3 trials

$$\binom{n=3}{k=3} = 1$$

$$P(3) = 1 * 0.5^3$$

$$\binom{n=3}{k=2} = 3$$

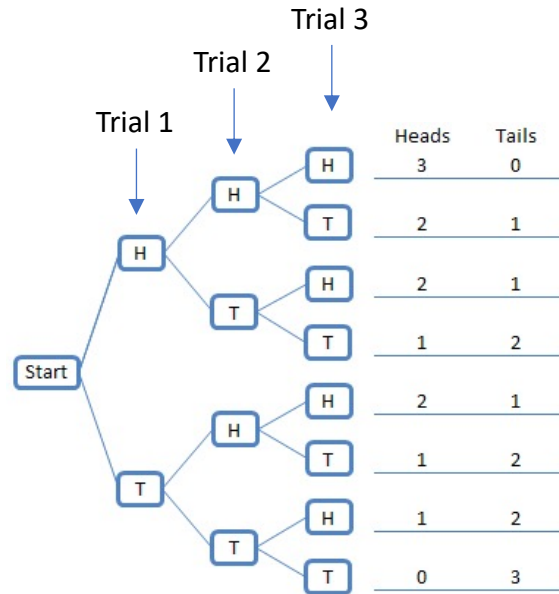
$$P(2) = 3 * 0.5^3$$

$$\binom{n=3}{k=1} = 3$$

$$P(1) = 3 * 0.5^3$$

$$\binom{n=3}{k=0} = 1$$

$$P(0) = 1 * 0.5^3$$



# Estimating probabilities

Number  $k$  heads in  
3 trials

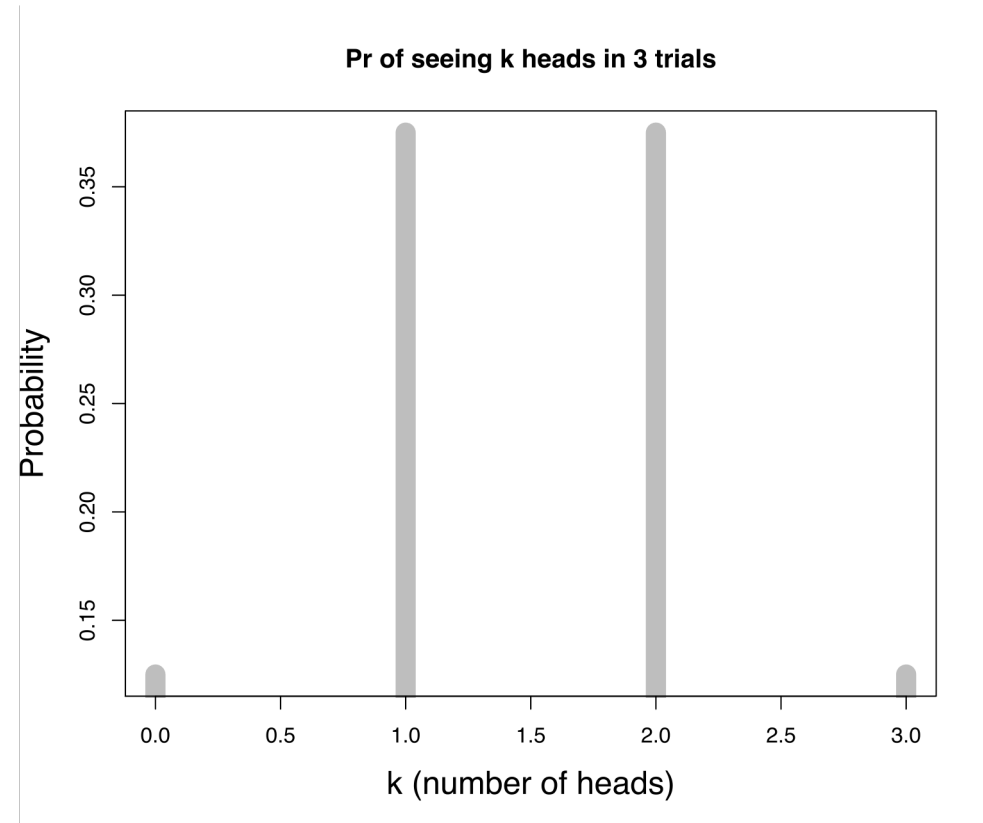
Probability of seeing  $k$  heads  
in 3 trials

$$\binom{n=3}{k=3} = 1 \quad P(3) = 1 * 0.5^3$$

$$\binom{n=3}{k=2} = 3 \quad P(2) = 3 * 0.5^3$$

$$\binom{n=3}{k=1} = 3 \quad P(1) = 3 * 0.5^3$$

$$\binom{n=3}{k=0} = 1 \quad P(0) = 1 * 0.5^3$$



# Binomial model ( and distribution)

Binomial model gives the probability of seeing  $k$  heads in  $n$  coin tosses (trials) given that probability of seeing a head in one coin toss is  $p$ .

Number of tosses in an experiment

Number of heads in an experiment

Number of ways to choose  $k$  heads

Probability of seeing  $k$  heads

Probability of seeing heads

Probability of seeing tails ( $n-k$ )

$$B(k|n, p) = \binom{n}{k} p^k (1-p)^{n-k}$$

$$\binom{n}{k} = \frac{n!}{k!(n-k)!}$$



- Coin is fair
- We toss the coin 3 times

# Binomial model

Number of heads in an experiment

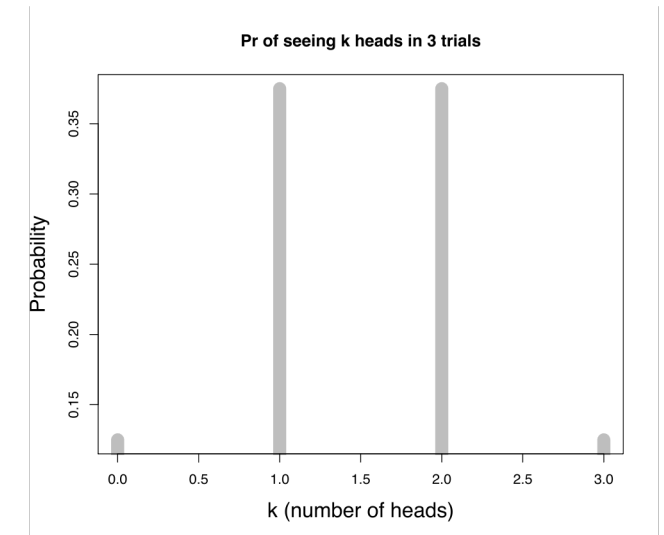

Number of tosses in an experiment

Number of ways to choose 2 heads in 3 tosses = 3

Probability of seeing heads

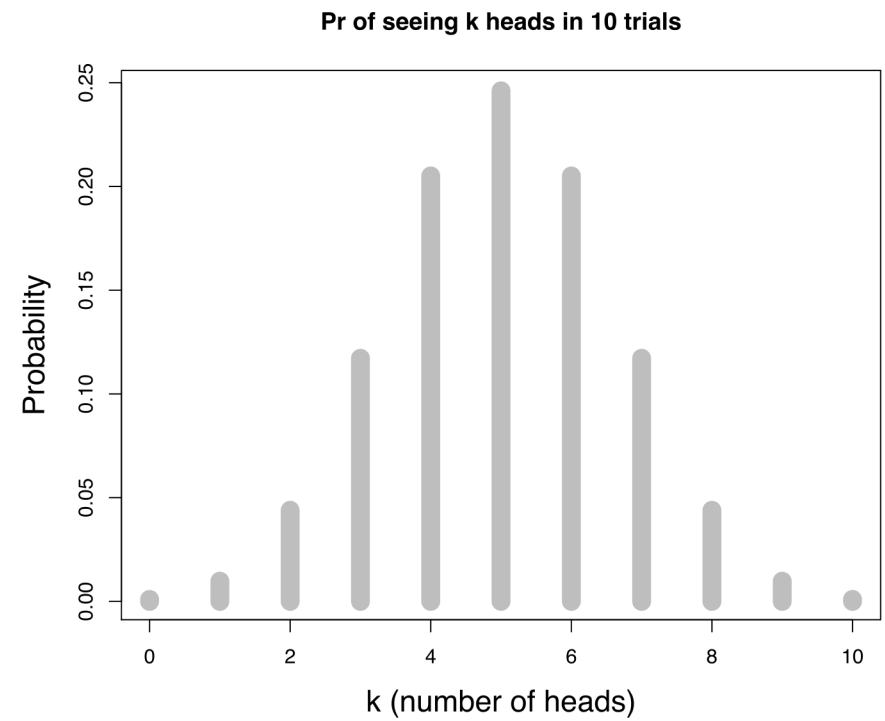
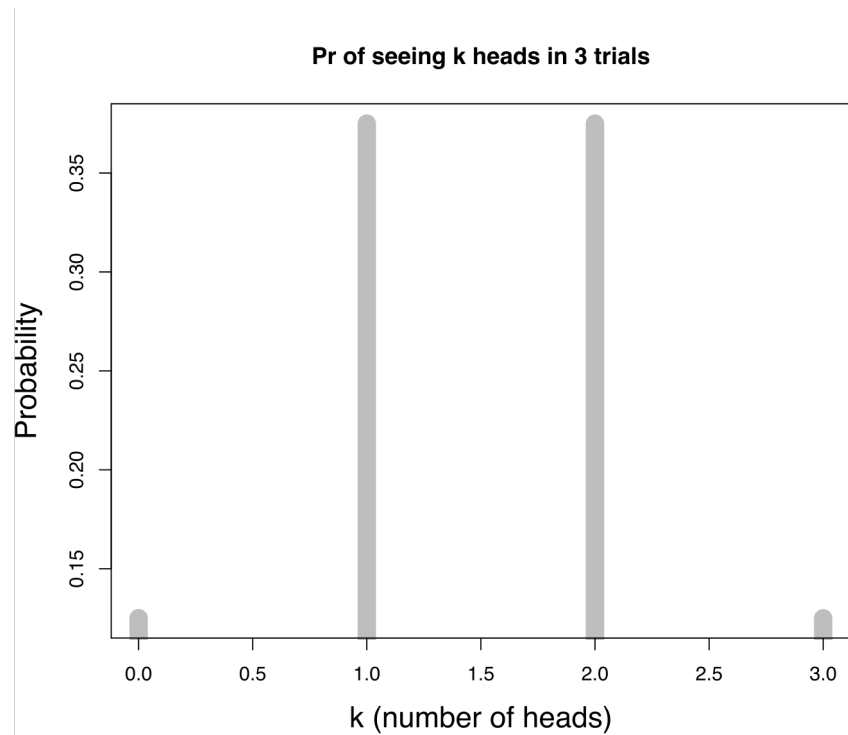
Probability of seeing 2 heads in 3 trials

Probability of seeing 2 heads and 1 tail =  $0.5^2 * 0.5$

$$B(2 | 3, 0.5) = \binom{3}{2} 0.5^2 (1 - 0.5)^{3-2} = 3 * 0.5^2 * 0.5 = 0.375$$




# Binomial model



# Likelihood function of Binomial distribution

In statistics, a likelihood function (often simply the likelihood) is a particular function of the parameter of a statistical model given data. Likelihood functions play a key role in statistical inference.

## Binomial model:

$P(k)$  of  $k$  heads given  $n$  trials and  $p$  of seeing a head in a trial

$$B(k|n, p) = \binom{n}{k} p^k (1 - p)^{n-k}$$

## Binomial Likelihood:

Given  $n$  and  $k$  infer  $p$  that maximizes the likelihood function

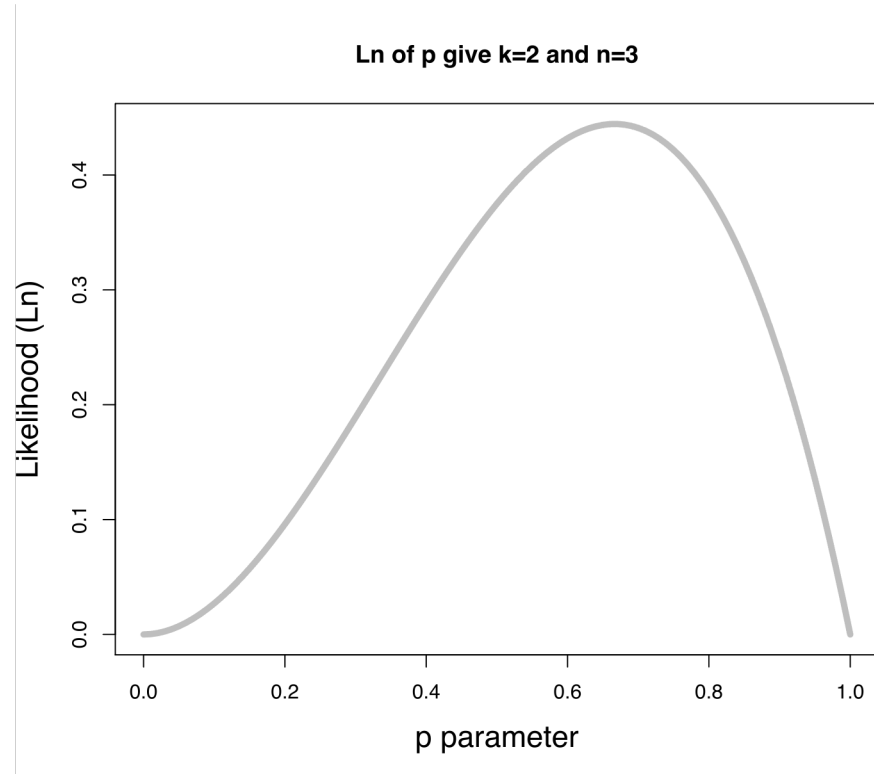
$$L(p | n = 3, k = 2) = \binom{n}{k} p^k (1 - p)^{n-k} = \binom{3}{2} p^2 (1 - p)^{3-2}$$



- Domain of  $p$  is a value between 0 and 1 (since  $p$  is a probability)
- Let's try all  $p$ 's to get a likelihood function
- Likelihood function is not a distribution

# Likelihood function of Binomial distribution

In statistics, a likelihood function (often simply the likelihood) is a particular function of the parameter of a statistical model given data. Likelihood functions play a key role in statistical inference.



## Binomial Likelihood:

Given  $n$  and  $k$  infer  $p$  that maximizes the likelihood function

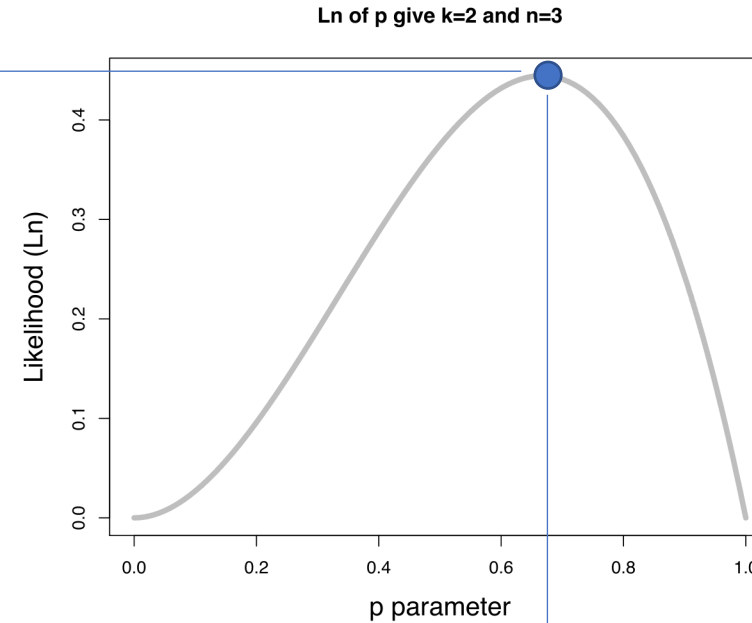
$$Ln(p | n = 3, k = 2) = \binom{n}{k} p^k (1 - p)^{n-k} = \binom{3}{2} p^2 (1 - p)^{3-2}$$



- Domain of  $p$  is a value between 0 and 1 (since  $p$  is a probability)
- Let's try all  $p$ 's to get a likelihood function
- Likelihood function is not a distribution

# Maximum Likelihood

Maximum value of the  
likelihood function  
 $Ln=0.44$



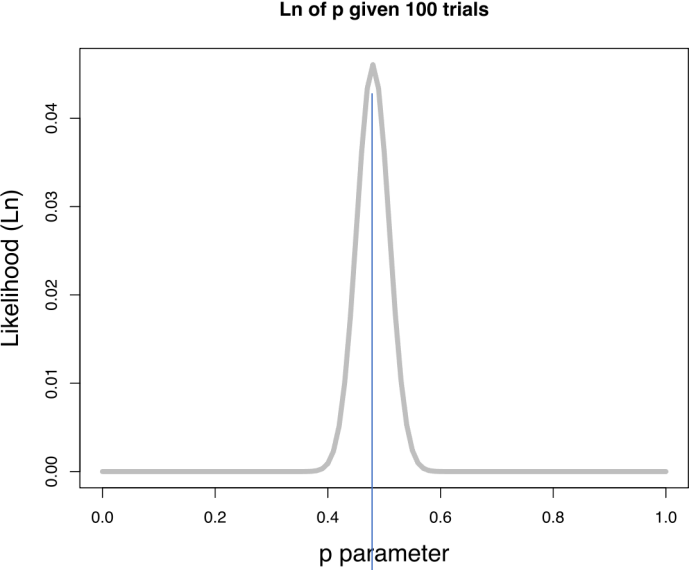
*Maximum likelihood estimate of the parameter*

$$Ln(\hat{p}) = \frac{k}{n} = \frac{2}{3} = 0.667$$

# Maximum Likelihood

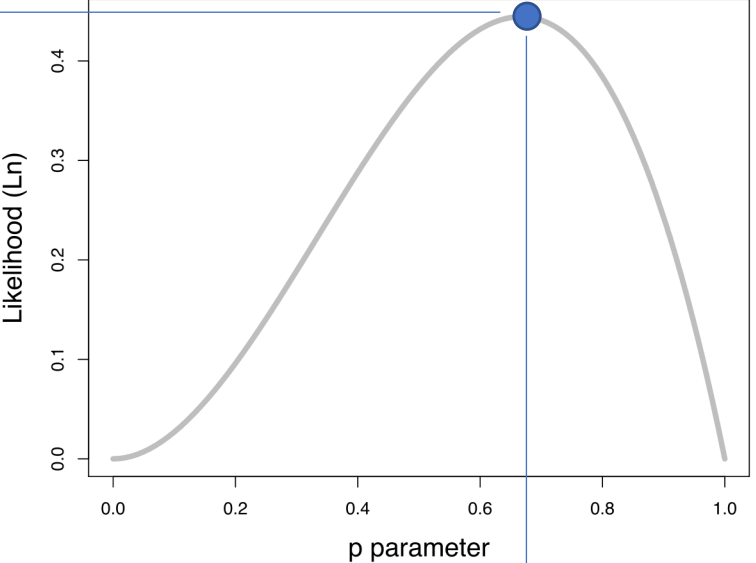


Maximum value of the likelihood function  
 $Ln=0.44$



$Ln(\hat{p}) = 0.48$

Ln of p give k=2 and n=3

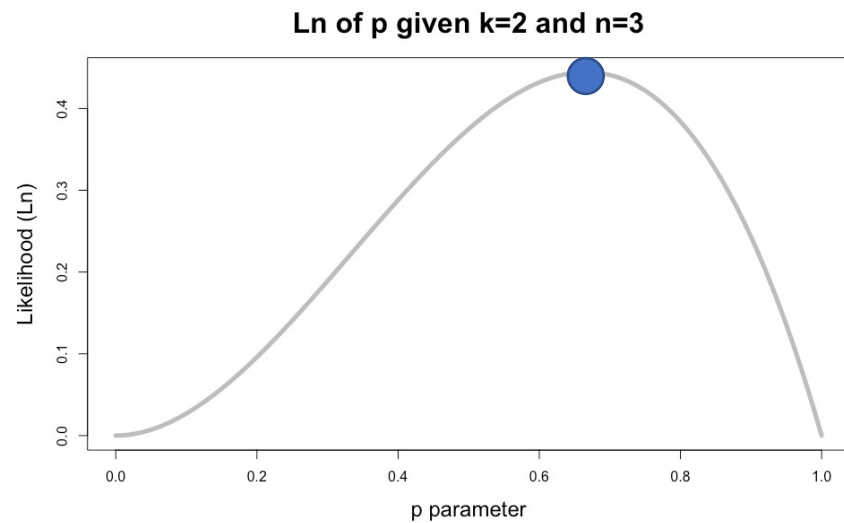


*Maximum likelihood estimate of the parameter*  
 $Ln(\hat{p}) = \frac{k}{n} = \frac{2}{3} = 0.667$

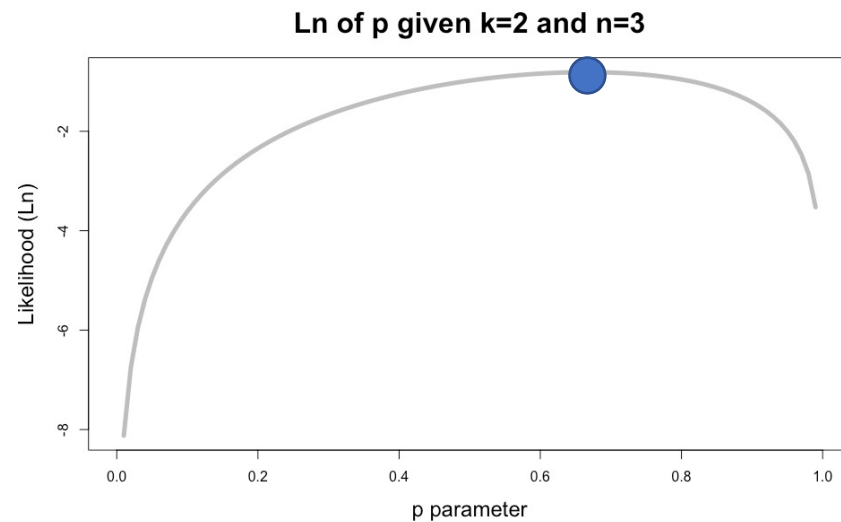
# Likelihood properties

- Likelihood is not a probability distribution!
- Log likelihood

likelihood



Log likelihood

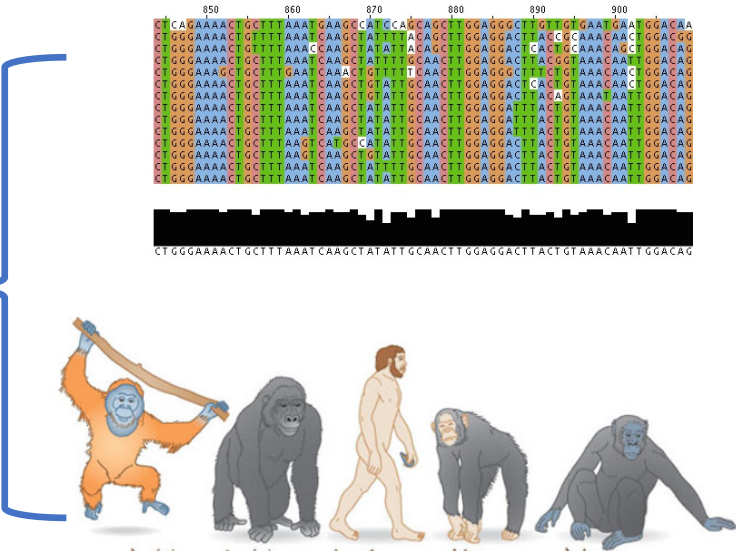


# Summary

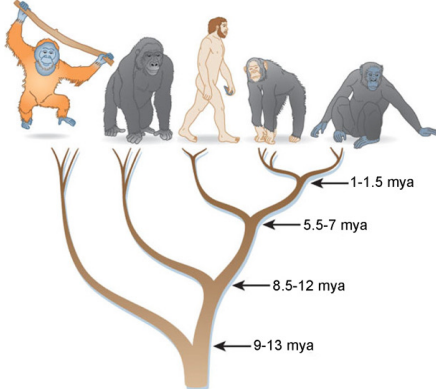
- Statistical phylogenetics is about modeling evolutionary process using probability distribution and stochastic processes
- Every measurement in this world is roughly speaking is a realization of some stochastic process
- In other words: every measurement is an instance that comes from some probability distribution (= model)
- Models are set of rules that describe how systems evolve
- In modeling data we need to come up with models that realistically describe our world
- **Maximum Likelihood method:** given that we observe an outcome and know the generating model, we can estimate the parameters of the process.

# Tomorrow's lecture

Observed data  
(DNA & Morphology)



inference





Let's calculate likelihood of some coin?

