

Model Selection

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Model selections vs model adequacy

Model Selection and Testing

General Introduction to Model selection

Comparing relative model fit with Bayes factors

Model selection of common substitution models for one locus

Comparing relative model fit with Bayes factors

Model selection of partition models

Comparing relative model fit with Bayes factors

Model averaging of substitution models

Reversible-jump MCMC over substitution models

Introduction to Posterior Prediction

Assessing the fit of Normal distributions to trait data

Assessing Phylogenetic Reliability Using RevBayes and P^3

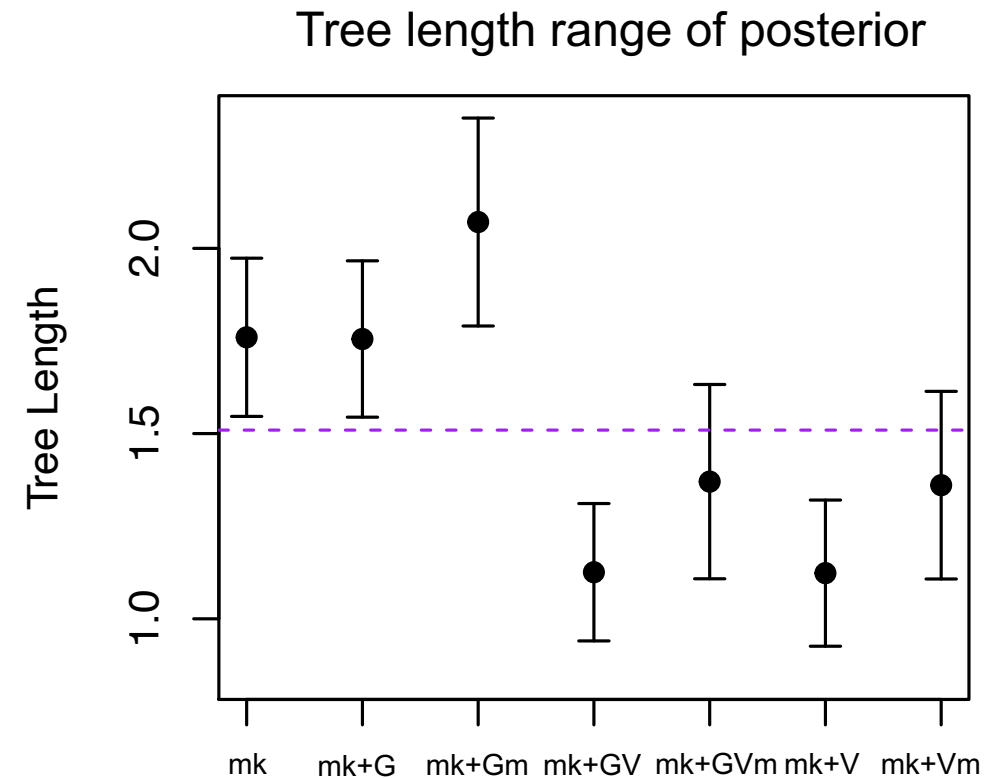
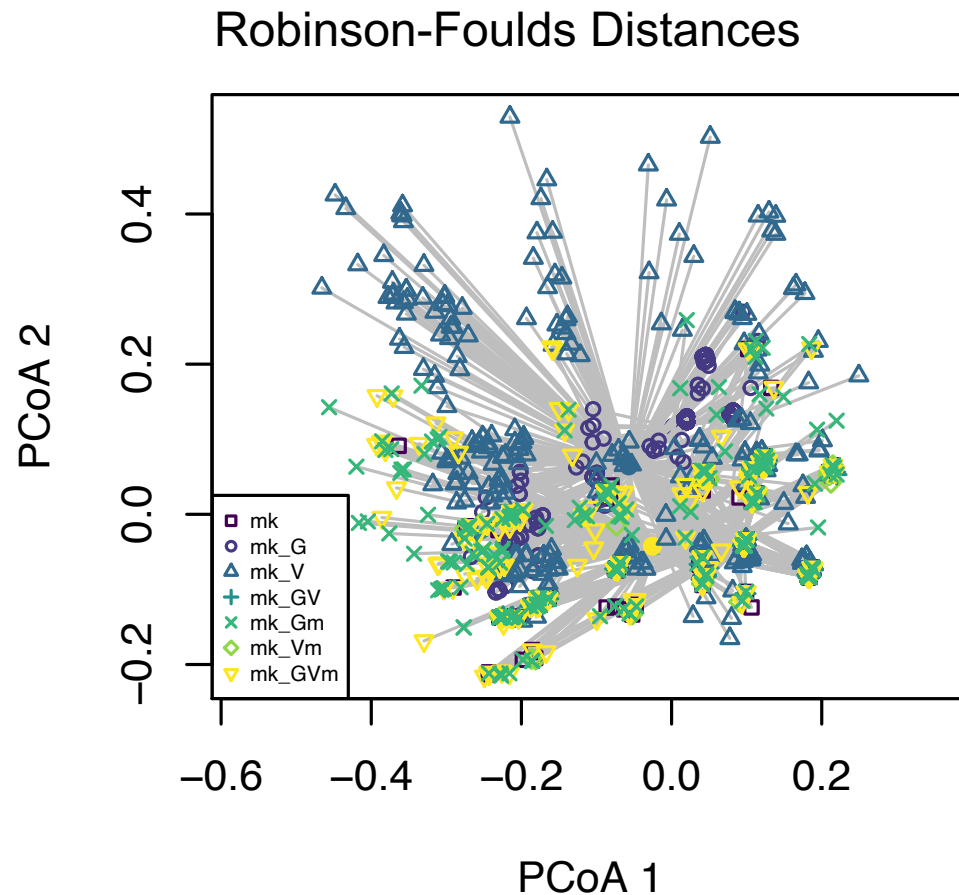
Model adequacy testing using posterior prediction (Data Version).

Assessing Phylogenetic Reliability Using RevBayes and P^3

Model adequacy testing using posterior prediction (Inference Version).

Revbayes [tutorials](#)

Impacts of substitution model on inferred parameters



(Mulvey et al in prep.)

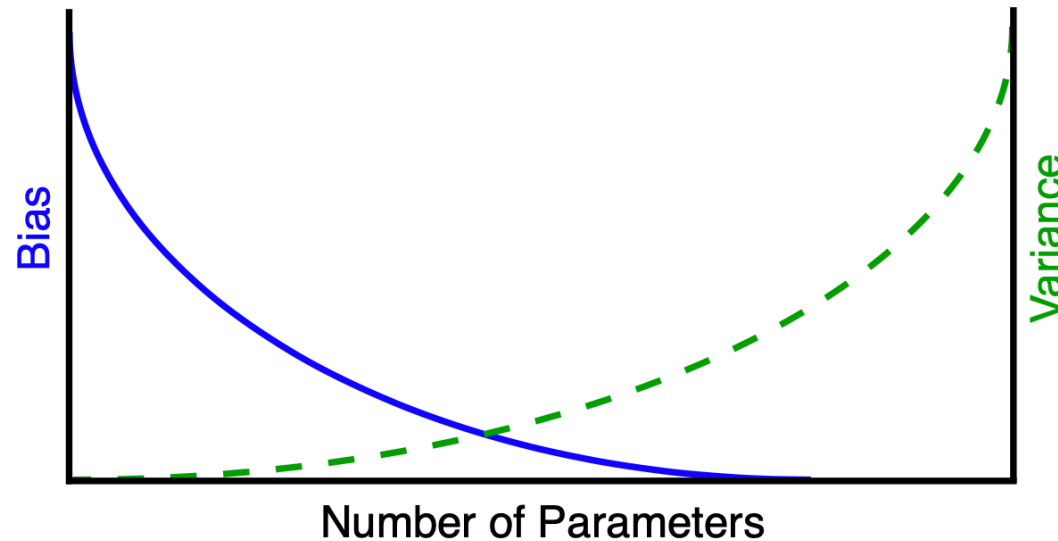
What does a good model look like?

To do statistical inference we need a model

What model should that be?

Our goal should be to have a model that is **complex enough** to capture the “important” variation in the data, but **not be more complex** than it needs to be

Too simple,
misinterpreting the
data



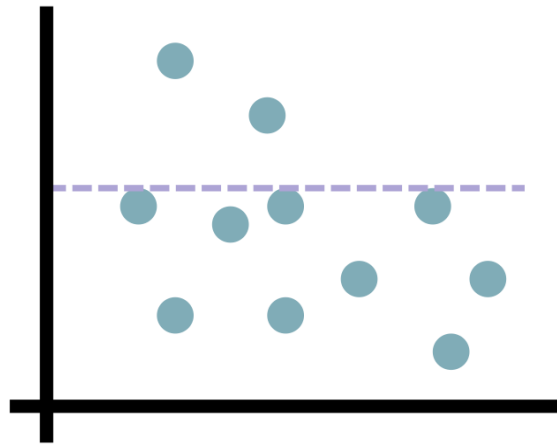
Too complicated,
not enough
information

What does a good model look like?

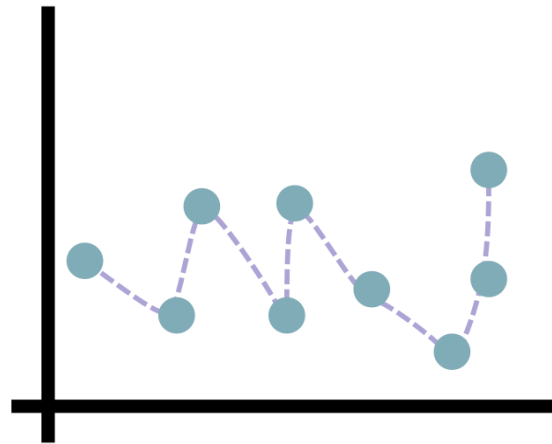
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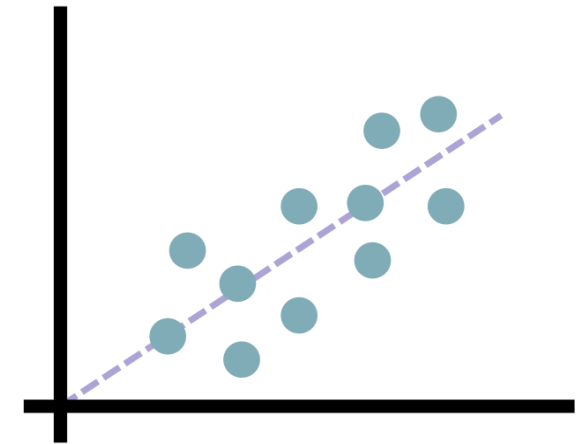
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Underfitting

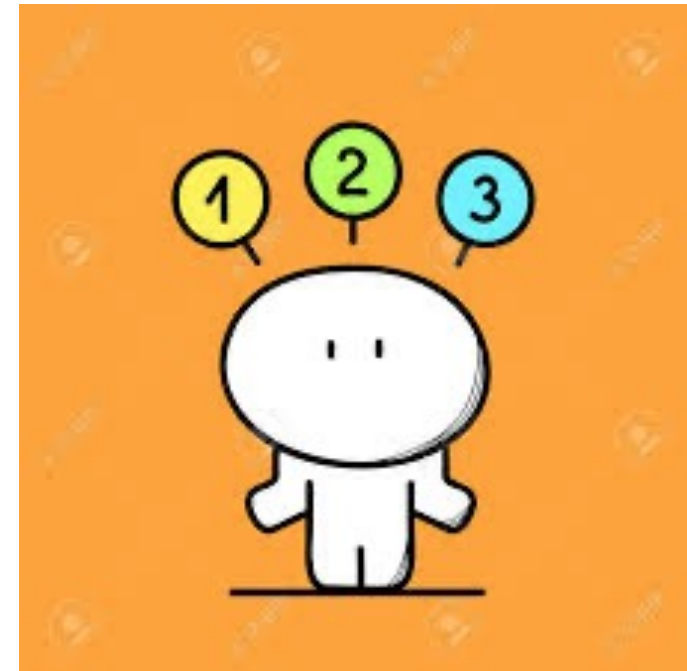


Overfitting



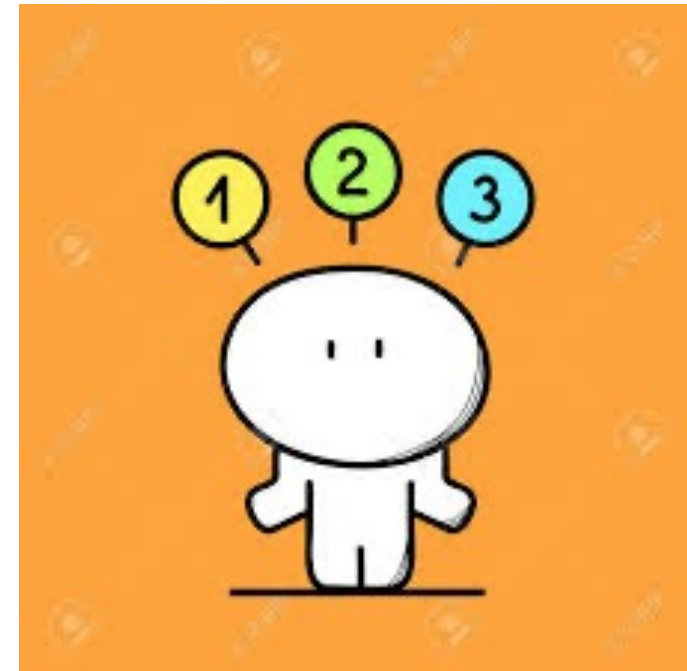
Proper fit

How to choose which model to use for morphological data?



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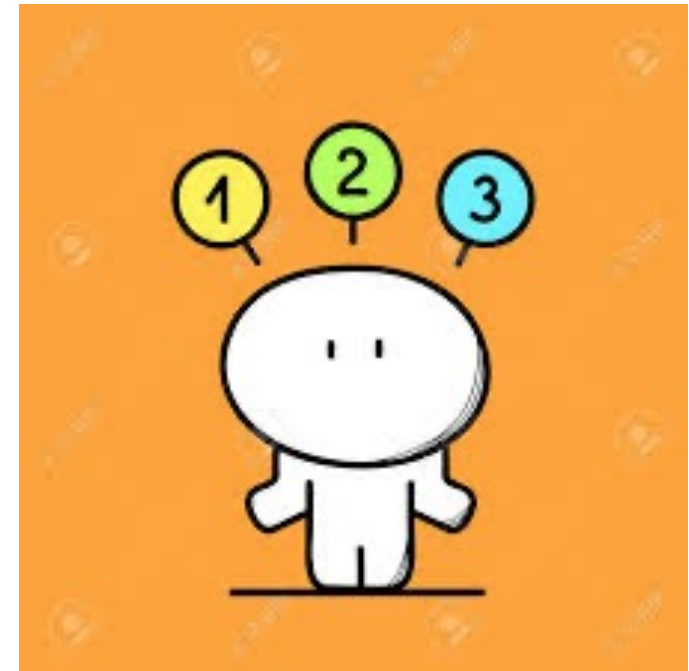
Guess



How to choose which model to use for morphological data?

Guess

What other people have done

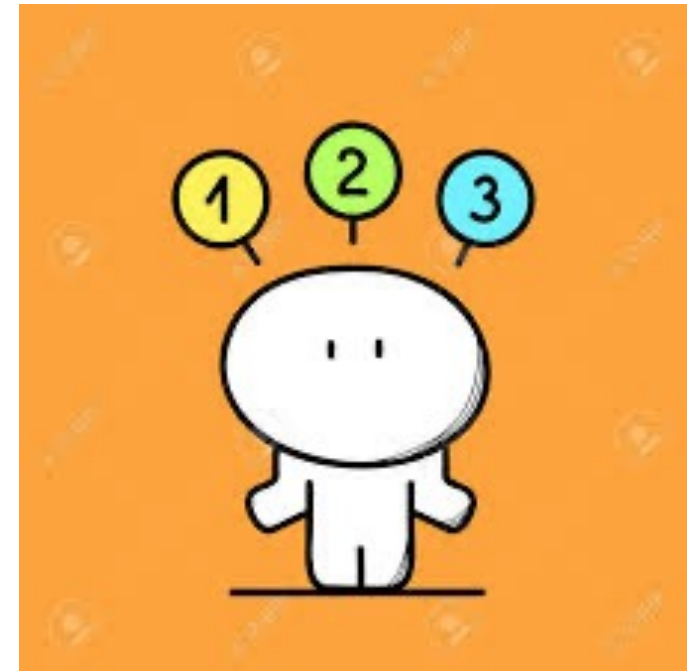


How to choose which model to use for morphological data?

Guess

What other people have done

AIC values



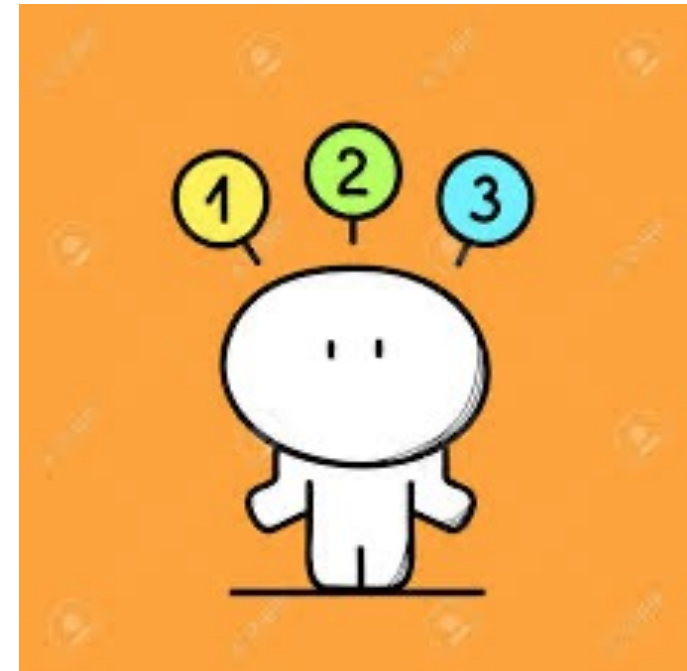
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What other people have done

AIC values

BIC Values



How to choose which model to use for morphological data?

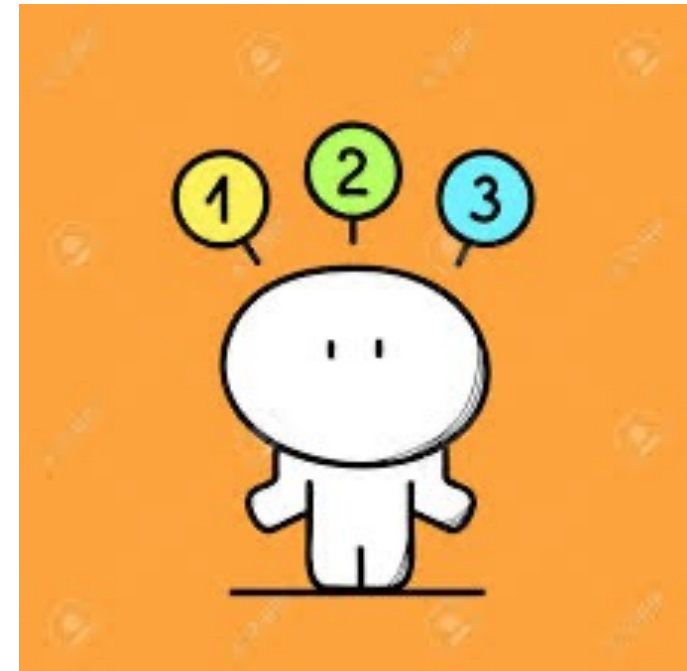
Guess

What other people have done

AIC values

BIC Values

Bayes factors :



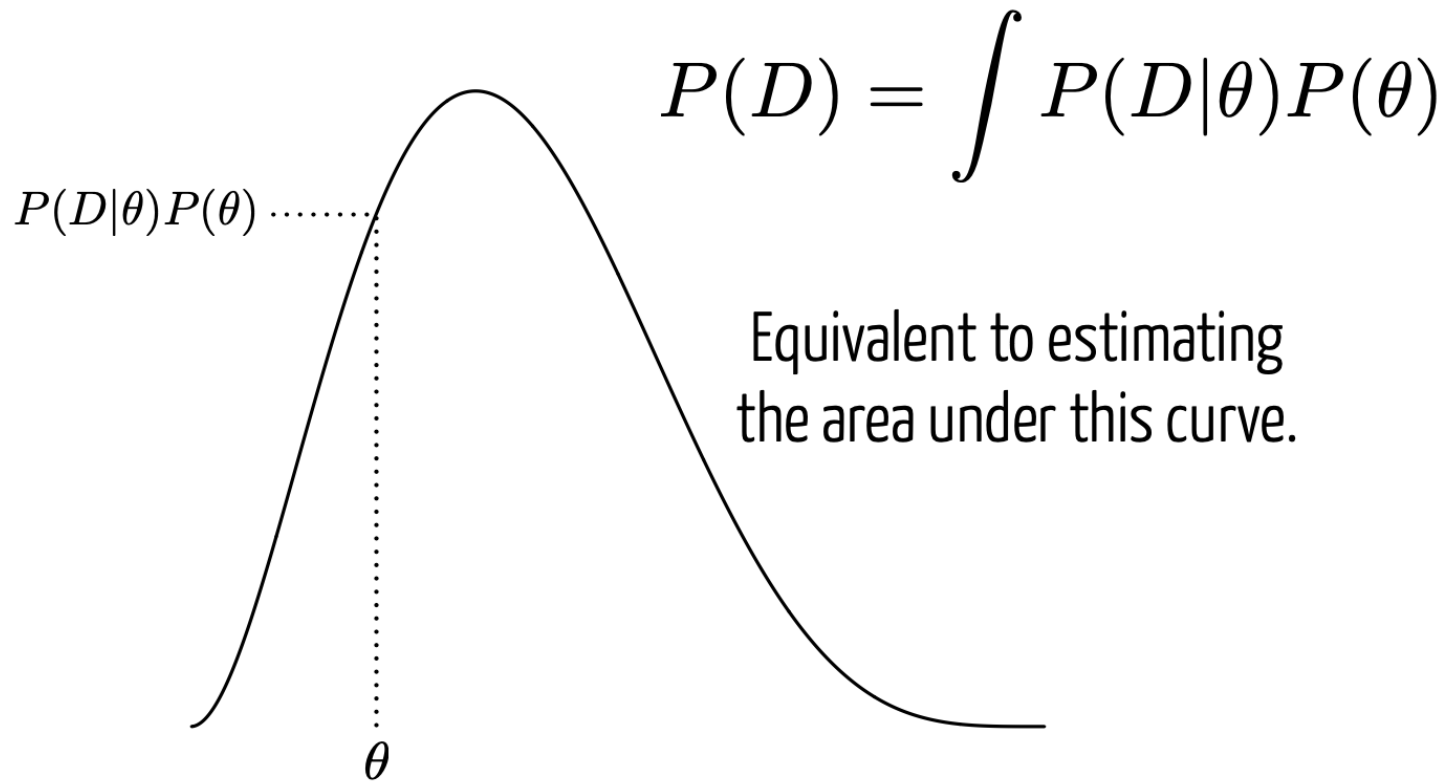
Bayes Theorem

$$P(\text{parameters} \mid \text{data}, \text{model}) =$$

↑
Posterior

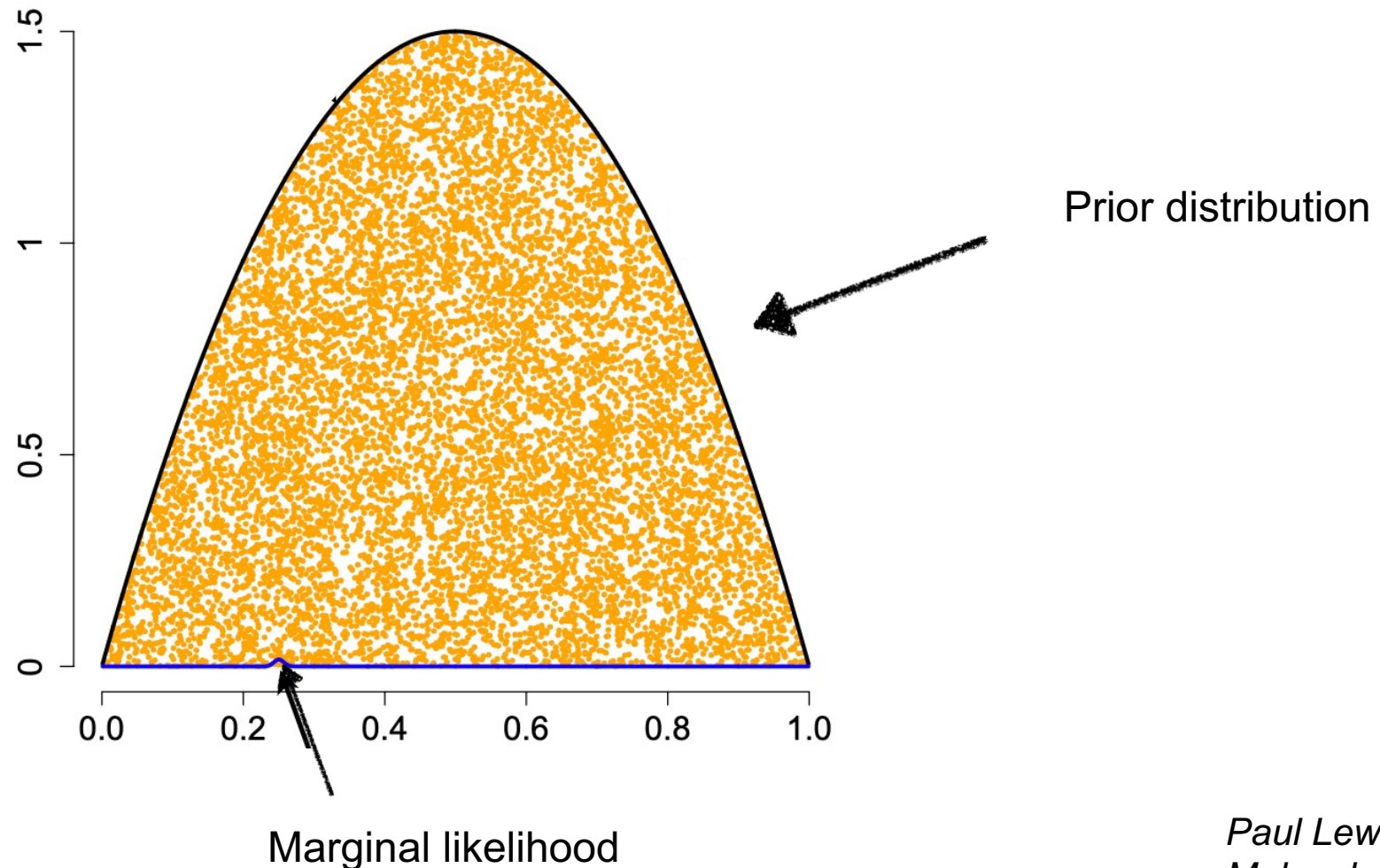
$$\frac{\begin{array}{c} \text{Likelihood} \\ \downarrow \\ P(\text{data} \mid \text{parameters}, \text{model}) \end{array} \begin{array}{c} \text{Priors} \\ \downarrow \\ P(\text{parameters} \mid \text{model}) \end{array}}{\begin{array}{c} \text{Marginal} \\ \text{probability} \uparrow \\ P(\text{data} \mid \text{model}) \end{array}}$$

What is the marginal likelihood.....



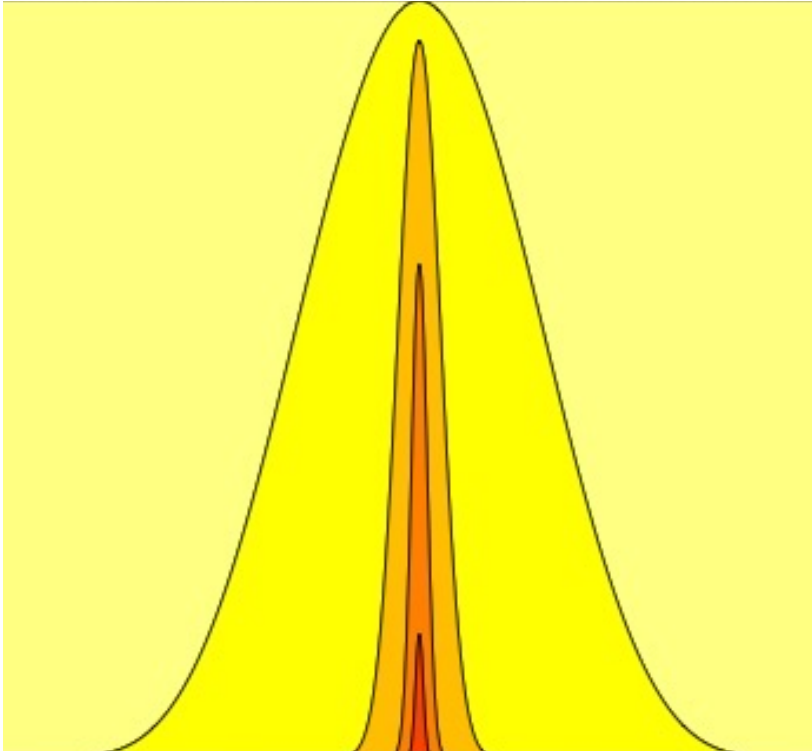
*Paul Lewis - Workshop on
Molecular Evolution 2016*

How can we estimate the marginal likelihood



*Paul Lewis - Workshop on
Molecular Evolution 2016*

Stepping Stone



Keep estimating smaller and smaller sections until you get down to the marginal likelihood

Paul Lewis - Phyloseminar

Exercise 2

Check which of the two models you chose for exercise 1 better than the other using a model selection approach.

Can you already predict which one you think might be better?