

Bayesian Tree Sampling

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CIBIV

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Outline

- 1 Bayes Theorem
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 - Its all about the die
 - Hats

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- 5 Phylogenetic Bayesian MCMC
 - In practice
 - Priors

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The difference

The Bayesian approach asks the right question in a hypothesis testing procedure, namely, “What is the probability that this hypothesis is true, given the data?” rather than the classical approach, which asks a question like, “Assuming that this hypothesis is true, what is the probability of the observed data?”

–Statistical Methods in Bioinformatics

Derivation

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from conditional probability. Also

$$\Pr(A \cap B) = \Pr(B \cap A) = \Pr(A|B) \Pr(B).$$

Therefore

$$\begin{aligned} \Pr(A|B) \Pr(B) &= \Pr(B|A) \Pr(A) \\ \Pr(A|B) &= \frac{\Pr(B|A) \Pr(A)}{\Pr(B)}. \end{aligned}$$

This is Bayes formula or theorem.

Bayes Theorem

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Bayes Theorem

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$$\underbrace{\Pr(A|B)}_{\text{Posterior Density}} \propto \underbrace{L(A, B)}_{\text{Likelihood}} \underbrace{\Pr(A)}_{\text{Prior}}$$

Bayes Theorem

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- Bayesian, flips the probability around.
- It is easy to include prior information which is often available.
- The Bayesian conditional probability is perhaps more intuitive.

Making formulas tangible

$$\Pr(T, M|D) \propto \Pr(D|T, M) \Pr(T, M)$$

- The likelihood is $L(T, D, M) = \Pr(D|T, M)$

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 - D is the DNA/Protein etc sequence data.
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- In words: The likelihood is the probability of the DNA data given the Tree and the model parameters.

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- In words: The likelihood is the probability of the DNA data given the Tree and the model parameters.
- The Prior is $\Pr(T, M)$ and indicates any information we already know. i.e. The root is not older than 10 million years.
- The Posterior density is $\Pr(T, M|D)$ the probability of the tree and model parameters given the sequence data.

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- Therefore MHMCMC must be used, this means it will take a lot of computer resources.
- The “answer” is not a tree, but a distribution of trees/states.
- It will always be slower than ML.

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Machine Output

1,2,1,0,1,0,-1,-2,-3,-2,-3,-4,-3,-2,-1,0,-1,0,1,2,1,2,3

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- The next item has a 50% chance that it will be a 4, and a 50% chance that it will be a 2.
- This is a Markov Chain.

Definition of a Markov Chain

Definition

A Markov Chain is a **chain** of randomly chosen values where the probability of the next value is entirely determined by the previous value.

Definition of a Markov Chain

Definition

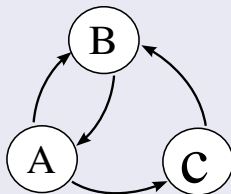
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Rough Math definition

$$\Pr(X_n | X_{n-1}, X_{n-2}, \dots) = \Pr(X_n | X_{n-1})$$

Markov Chain Graph

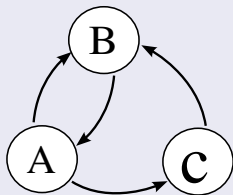
State Graph



- Simple Markov Chains can be represented as a graph.

Markov Chain Graph

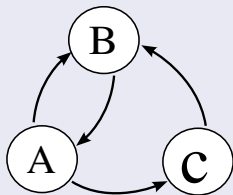
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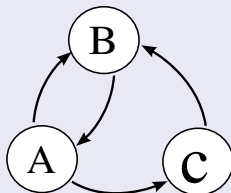
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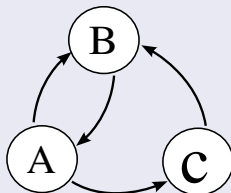
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Markov Chain Graph

State Graph



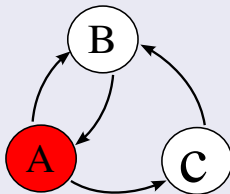
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- For clarity, when transitions are equiprobable we omit the transition probabilities.

definition

Markov Chain Graph

Example

State Graph



Output

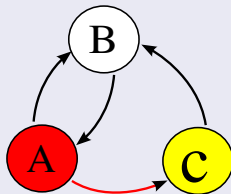
A

definition

Markov Chain Graph

Example

State Graph



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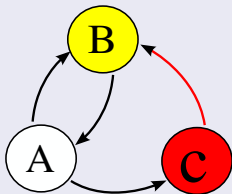
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definition

Markov Chain Graph

Example

State Graph



Output

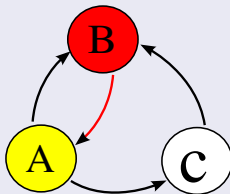
A C B

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Markov Chain Graph

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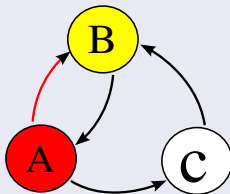
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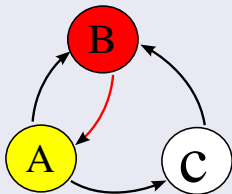
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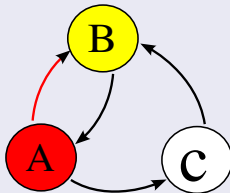
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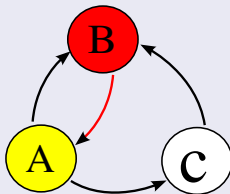
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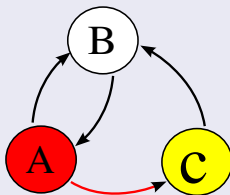
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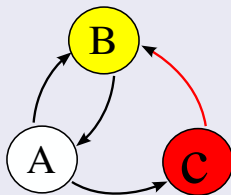
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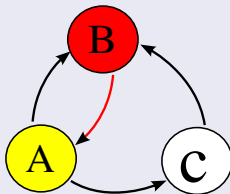
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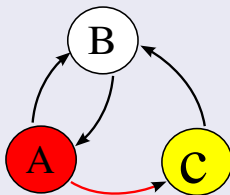
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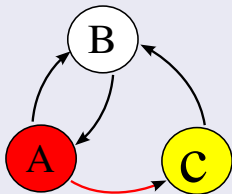
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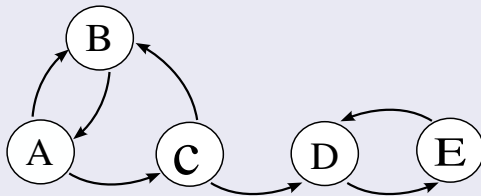
- Note that the **states** can be anything. ie different trees

Properties

Extra Markov Chain Properties

Irreducibility

Reducible state diagram

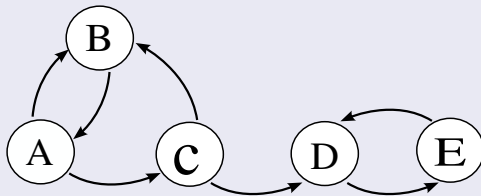


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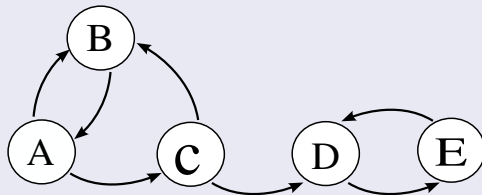
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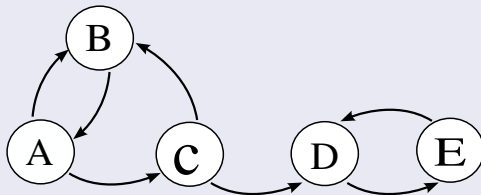
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A Markov Chain is **Irreducible** if and only if the chain can get from **any** possible state to **any** other possible state eventually.

Extra Markov Chain Properties

Irreducibility

Reducible state diagram



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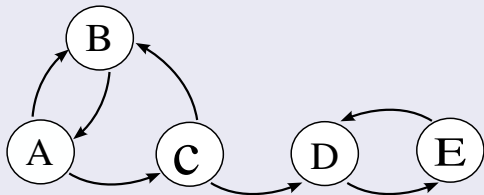
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- The above state diagram is **NOT** irreducible.

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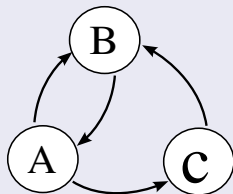
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A Markov Chain is **Irreducible** if and only if the chain can get from **any** possible state to **any** other possible state eventually.

- The above state diagram is **NOT** irreducible.
- Adding a transition from $D \rightarrow C$ it would make this irreducible

Extra Markov Chain Properties

Reversibility



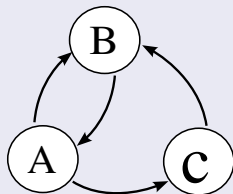
Is this output reversed?

C A B C A B A B A B C

- Note that there is no $C \rightarrow B$ transition or $C \rightarrow A$ transition.

Extra Markov Chain Properties

Reversibility



Is this output reversed?

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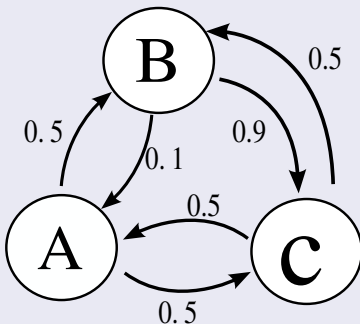
- Note that there is no $C \rightarrow B$ transition or $C \rightarrow A$ transition.
- Therefore we can tell that this output sequence is reversed.

Properties

Extra Markov Chain Properties

Reversibility

Tricky Example



Is this output reversed?

ABCABCBCBCBCABCBCABA

Extra Markov Chain Properties

Reversibility

Is this output reversed?

A B C A B C B C B C B C A B C B C A B A

- The transition $B \rightarrow A$ is much less likely than $B \rightarrow C$ in the forward direction.

Extra Markov Chain Properties

Reversibility

Is this output reversed?

A B C A B C B C B C B C A B C B C A B A

- The transition $B \rightarrow A$ is much less likely than $B \rightarrow C$ in the forward direction.
- In this example there are 7 $B \rightarrow C$ transitions and only 1 $B \rightarrow A$ transition in the forward direction.

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- Conversely there are 4 $B \rightarrow C$ transitions and 4 $B \rightarrow A$ transitions in the reverse direction.
- It seems we can guess that this output is not reversed.
- But we stick to simple definitions for this course.

Extra Markov Chain Properties

Reversibility

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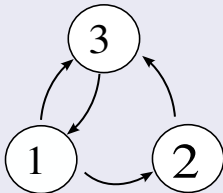
A Markov Chain is reversible if we cannot detect whether or not the chain is running in “reverse”. That is the output is statistically identical in both directions.

Properties

Extra Markov Chain Properties

Aperiodic

Periodic-Aperiodic

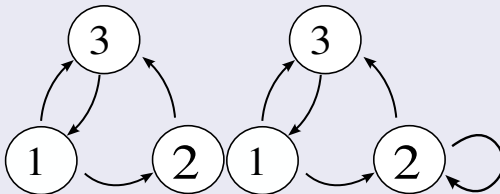


Properties

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Aperiodic

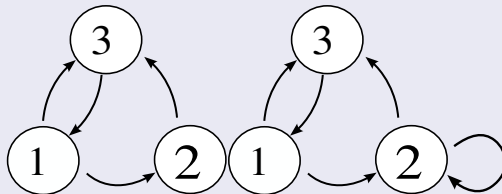
Periodic-Aperiodic



Extra Markov Chain Properties

Aperiodic

Periodic-Aperiodic



Definition

A Markov Chain is periodic if there is some fixed “cycle” of states, and it is aperiodic otherwise.

Extra Markov Chain Properties

Why do we care?

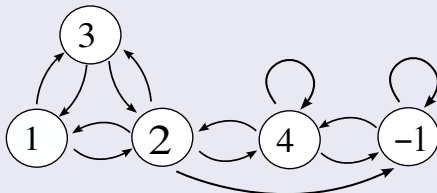
Extra Markov Chain Properties

Why do we care?

- If a MCMC chain has these 3 properties (reversible, irreducible and aperiodic), then it is also ergodic.

Extra Markov Chain Properties

Stationary distribution



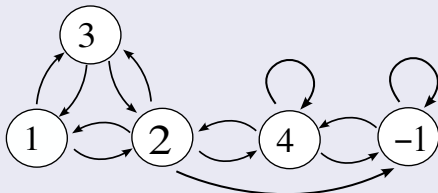
output

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- We can calculate statistics on the output, like mean and standard deviation. Also we can plot histograms etc.

Extra Markov Chain Properties

Stationary distribution



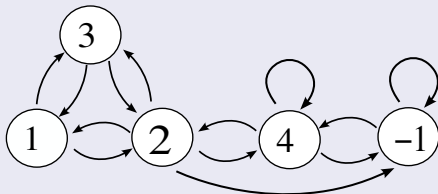
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- Consider the **distribution** of the output.
- What about the start state. That is if the chain is started in state 1, will the distribution be different from starting in 2.

Extra Markov Chain Properties

Ergodic

Definition

If we can start from **any** state, and if we take samples for long enough, and we end up with the same distribution, that distribution is the stationary distribution of the Markov Chain, and the Markov Chain is said to be **ergodic**

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- Usually the symbol π denotes the stationary distribution.
- Note that we have not said anything about how many **samples** we need to get an accurate distribution.

Outline

- 1 Bayes Theorem
 - Bayes Theorem
- 2 Markov Chains
 - definition
 - Properties
- 3 **MHMCMC**
 - Algorithm
 - Examples
- 4 What is long enough
 - Its all about the die
 - Hats
- 5 Phylogenetic Bayesian MCMC
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Metropolis Hastings MCMC

Algorithm

- Start in state X_n

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- If we accept, then $X_{n+1} = X'$, otherwise $X_{n+1} = X_n$.

Algorithm

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The Key Idea

The stationary distribution **is the posterior distribution** of interest. That is the MHMCMC chain is sampling the Bayesian posterior distribution.

Example

- Start with tree $T = (a, b|c, d)$.

Output

$(a, b|c, d)$

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- The new state is $T = (a, d|b, c)$
- We continue $T' = (a, c|b, d)$ ($c \rightleftharpoons d$), and accept.

Output

$(a, b|c, d)$ $(a, b|c, d)$ $(a, d|b, c)$ $(a, c|b, d)$

Die example

Wiki Formula

$$\Pr(k|i, s) = \frac{1}{s^i} \sum_{n=0}^{\lfloor \frac{k-i}{s} \rfloor} (-1)^n \binom{i}{n} \binom{k - sn - 1}{i - 1}$$

Die MHMCMC

- Formula looks too complicated!

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- Use a simple MHMCMC instead.
- Just pick one die at random and re-throw.
- This is reversible and the acceptance ratio is 1. i.e we always accept.

Examples

Die example

3 die

1 1 1

Output

3

Examples

Die example

3 die

1	1	1
4	1	1

Output

3 6

Examples

Die example

3 die

1	1	1
4	1	1
4	1	6

Output

3 6 11

Die example

3 die

1	1	1
4	1	1
4	1	6
2	1	6

Output

3 6 11 9

Die example

3 die

1	1	1
4	1	1
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2	1	6
3	1	6

Output

3 6 11 9 10

Die example

3 die

1	1	1
4	1	1
4	1	6
2	1	6
3	1	6
3	1	4

Output

3 6 11 9 10 8

Die example

3 die

1	1	1
4	1	1
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2	1	6
3	1	6
3	1	4
3	5	4

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3 6 11 9 10 8 12

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- By changing just one dice at each step, the sum can never change by more than 5 from step to step.

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Its all about the die

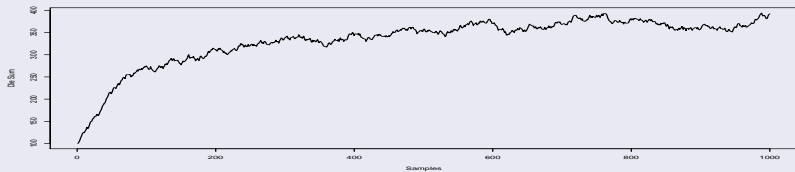
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- In this case we get to equilibrium in just a single step but must generate 100 random numbers per step.

Its all about the die

More Die

100 die, rolling 1 dice per step



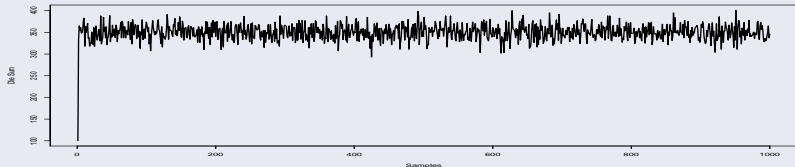
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More Die

100 die, rolling 1 dice per step



100 die, rolling all per step



Its all about the die

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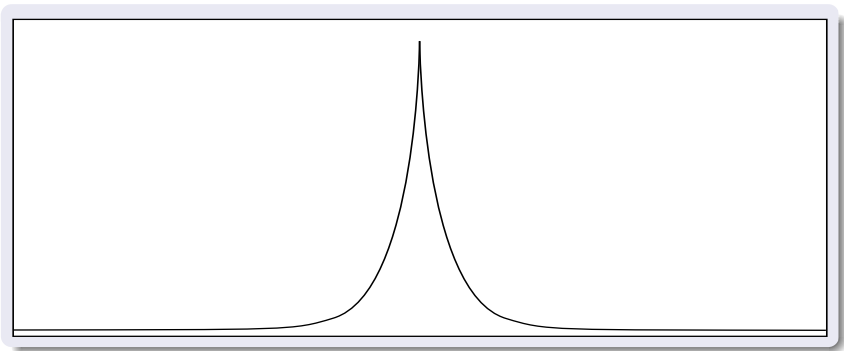
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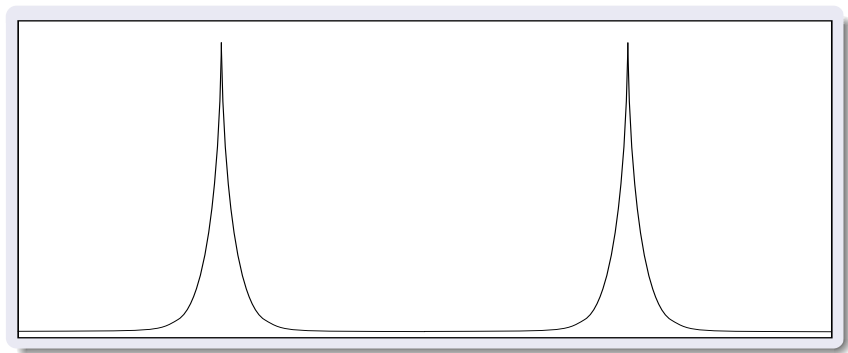
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- Due to the correlations between samples we don't really need every sample from the MCMC chain and instead only collect every 100'th sample or so.
- Performance should be measured in the number of effective samples per CPU cycle.

Hats

Witch's Hat



Witch's Hat



- Consider all non tree like signals.
- Recombination, Horizontal Gene Transfer and other effects could contribute to a lot of witch's hats.

Hats

Key Points for simple analysis

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- Multiple runs from random starting locations

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In practice

Posterior

$$\Pr(T, M|D) \propto \Pr(D|T, M) \Pr(T, M)$$

- The likelihood is $L(T, D, M) = \Pr(D|T, M)$
- T is the tree.
- D is the DNA/Protein etc sequence data.
- M is the model parameters, like GTR.

Warning

Trees Make Life Difficult

In practice

Moves and why you care about irreducibility

- Many programs have a huge set of options.

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- Examples of real output.

In practice

Aside: Hot and Cold chains

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- Only collect samples from the cold chain. ie the only chain with the correct distribution.
- The idea is that we won't get stuck.
- Generally not as effective as just developing some better moves.

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- Priors should be considered with respect to the hypothesis that will be tested.

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- Even if the max root height is 100 expected substitutions per site, the posterior can now be normalized.

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- Check your priors!

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 - Generally slower than ML. (bootstrapped)
 - Support values are easier to interpret.
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