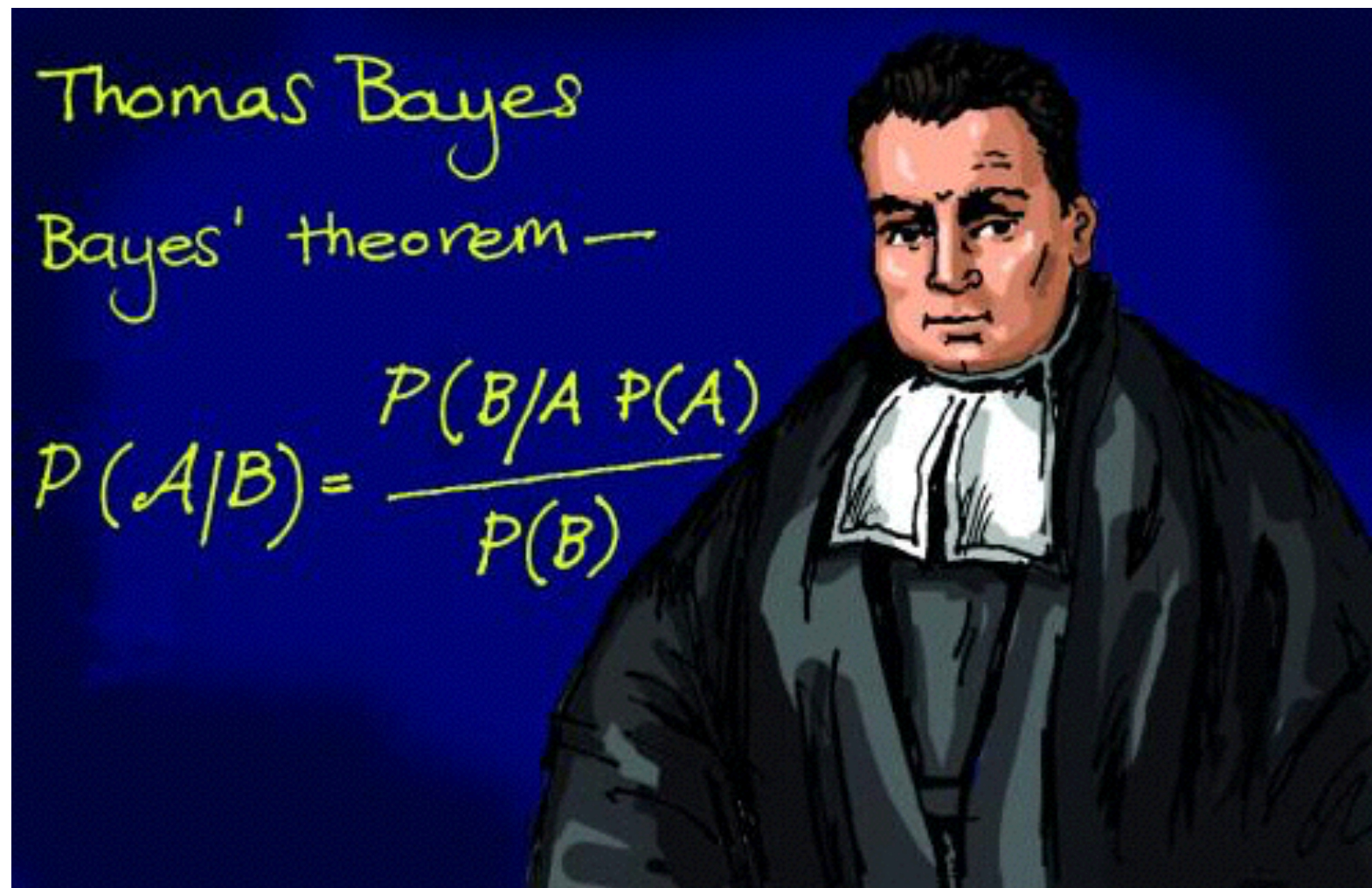


A Brief Overview of Bayesian Inference



Some Facts About Probability

$$P(A \cap B) = P(A|B)P(B) = P(B|A)P(A)$$

$$P(A|B)P(B) = P(B|A)P(A)$$

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

Bayes Theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$



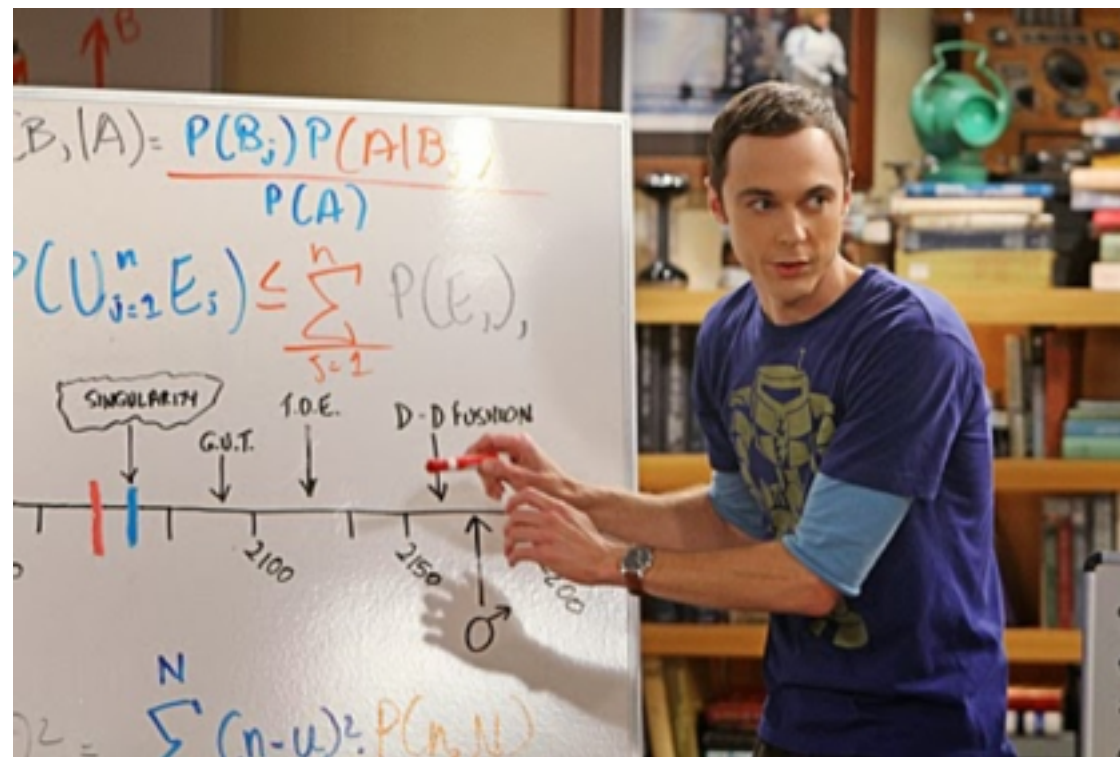
T. Bayes.

Bayes Theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$



T. Bayes.

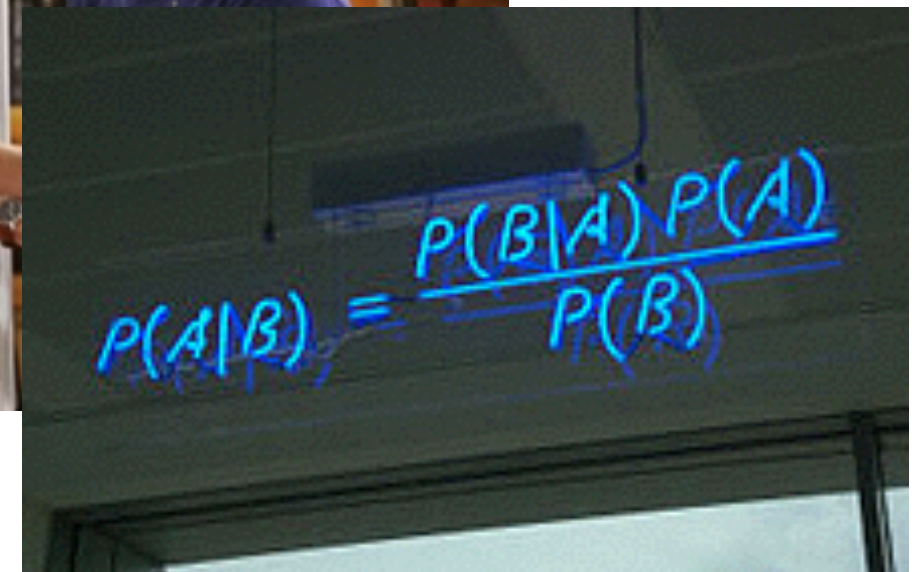
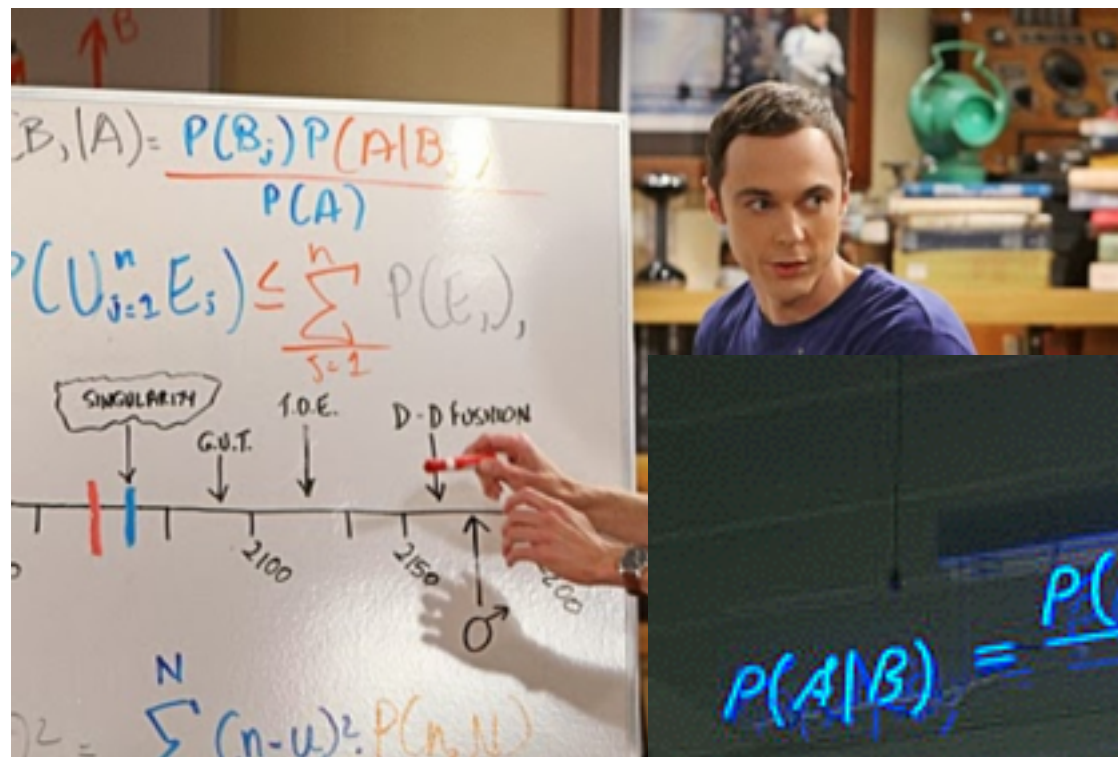


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T. Bayes.

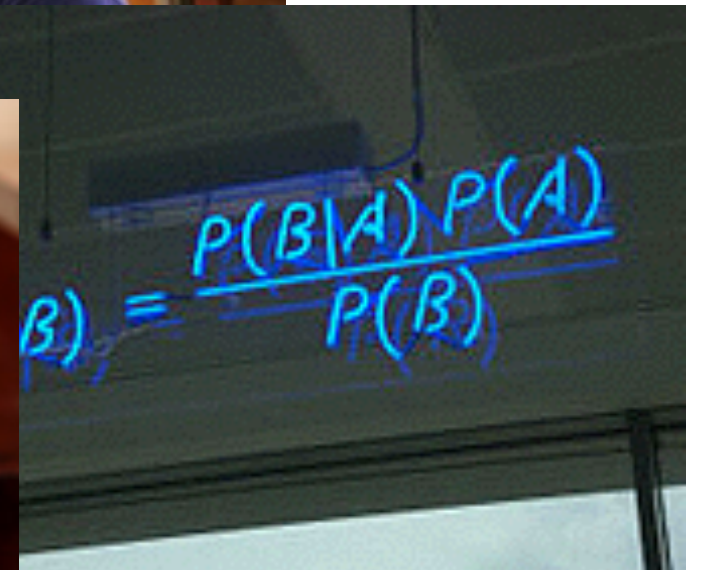
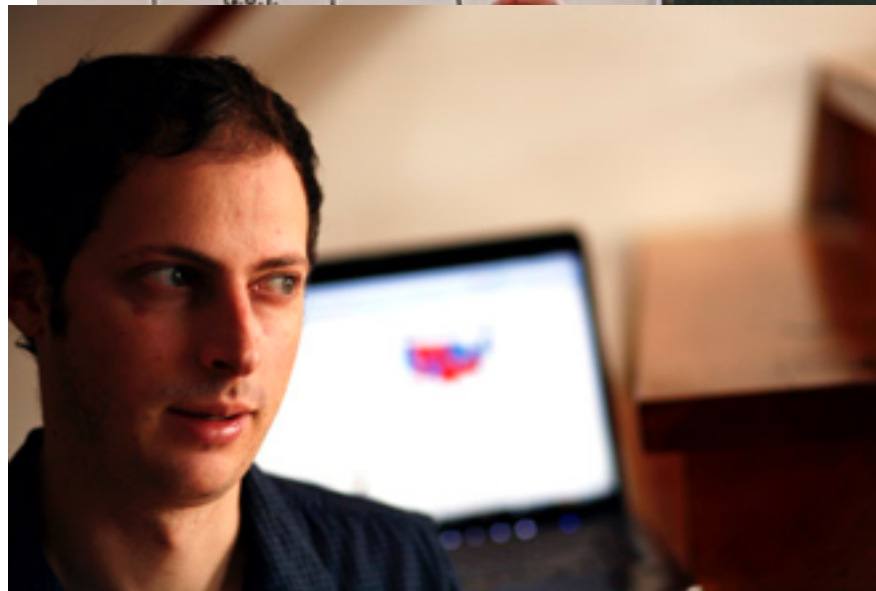
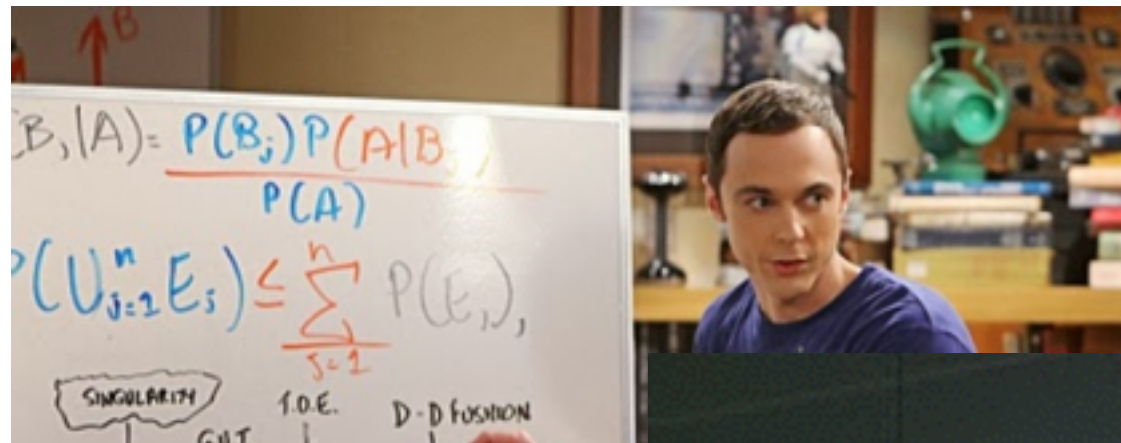


Bayes Theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$



T. Bayes.

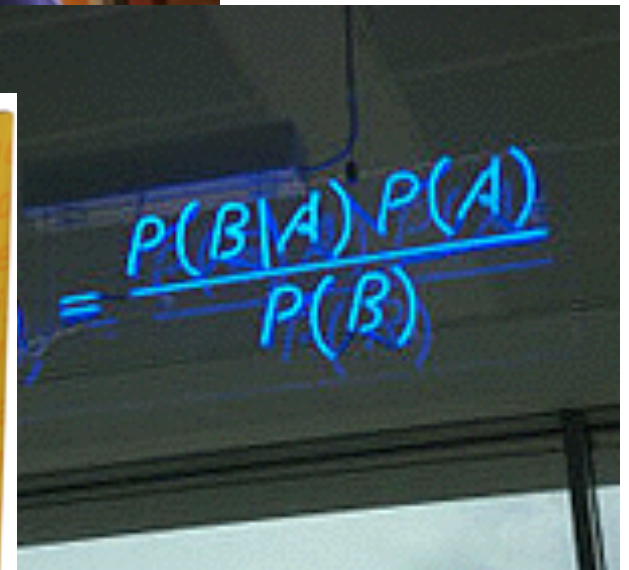
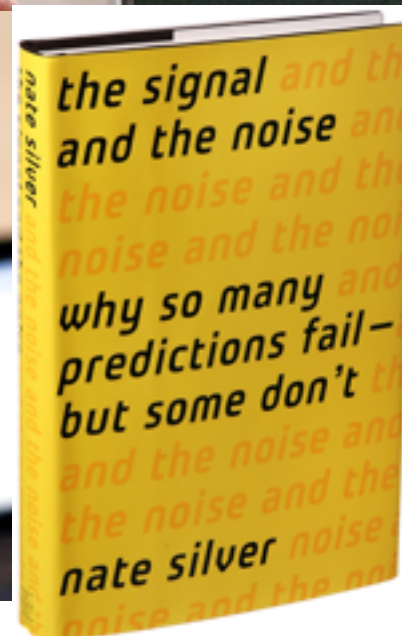
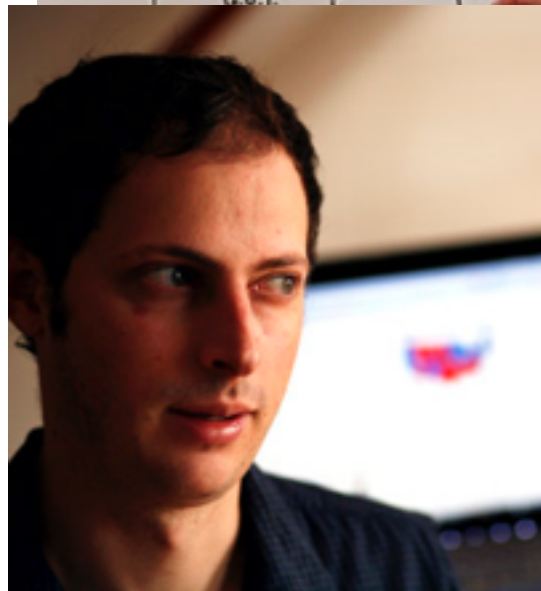
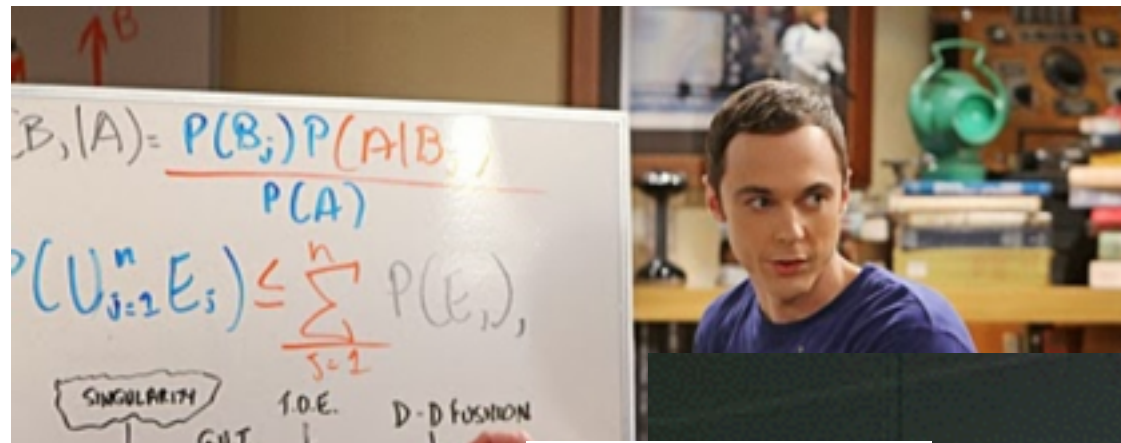


Bayes Theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$



T. Bayes.



Bayes Theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$



T. Bayes.

Rev. Thomas Bayes
1701-1761

“An Essay towards solving a
Problem in the Doctrine of Chances”
published in 1763 (Richard Price)

Binomial with a uniform prior on p

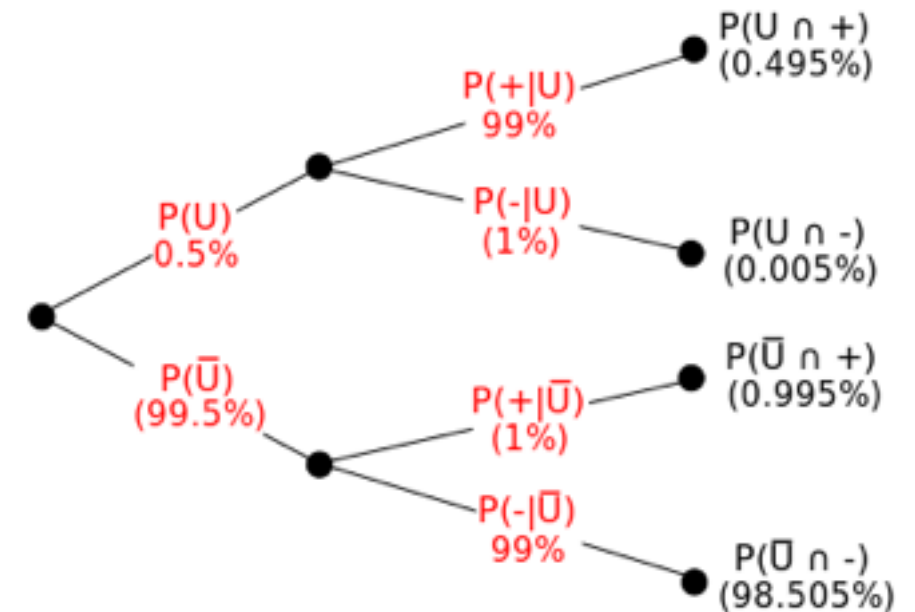
Bayes Theorem

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

Also used with frequentist probabilities!

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

Suppose a drug test is 99% sensitive and 99% specific. That is, the test will produce 99% true positive results for drug users and 99% true negative results for non-drug users. Suppose that 0.5% of people are users of the drug. If a randomly selected individual tests positive, what is the probability that he is a user?



$$\begin{aligned}
 P(\text{User} \mid +) &= \frac{P(+ \mid \text{User})P(\text{User})}{P(+ \mid \text{User})P(\text{User}) + P(+ \mid \text{Non-user})P(\text{Non-user})} \\
 &= \frac{0.99 \times 0.005}{0.99 \times 0.005 + 0.01 \times 0.995} \\
 &\approx 33.2\%
 \end{aligned}$$

Bayes Theorem

Posterior

Likelihood

Prior

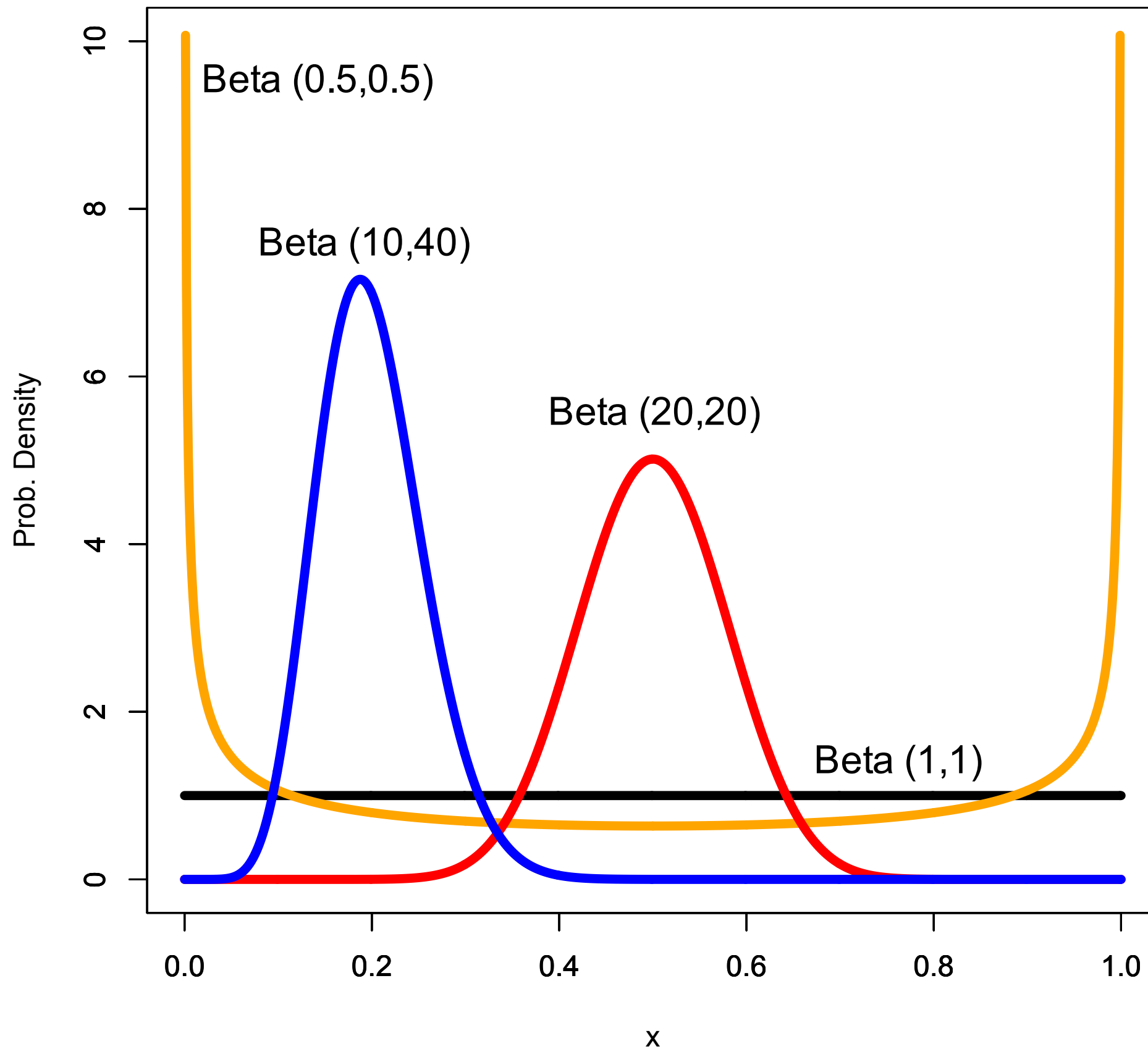
$$P(H|D) = \frac{P(D|H)P(H)}{P(D)}$$

Normalizing Constant
(Marginal Likelihood)

The diagram illustrates the components of Bayes' Theorem. The equation $P(H|D) = \frac{P(D|H)P(H)}{P(D)}$ is centered. A red arrow points from the word 'Posterior' to the term $P(H|D)$. A blue arrow points from the word 'Likelihood' to the term $P(D|H)$. A green arrow points from the word 'Prior' to the term $P(H)$. An orange arrow points from the text 'Normalizing Constant (Marginal Likelihood)' to the term $P(D)$ in the denominator.

Beta distributions are a flexible class of possible priors for continuous numbers between 0 and 1.

Beta Distributions



Odds Ratios

$$\frac{P(H_1|D)}{P(H_2|D)} = \frac{\frac{P(H_1)P(D|H_1)}{P(D)}}{\frac{P(H_2)P(D|H_2)}{P(D)}}$$

Odds Ratios

$$\frac{P(H_1|D)}{P(H_2|D)} = \frac{\frac{P(H_1)P(D|H_1)}{P(D)}}{\frac{P(H_2)P(D|H_2)}{P(D)}}$$

Odds Ratios

$$\frac{P(H_1|D)}{P(H_2|D)} = \frac{P(H_1)P(D|H_1)}{P(H_2)P(D|H_2)}$$

Odds Ratios

Posterior Odds

Prior Odds

Bayes Factor

$$\frac{P(H_1|D)}{P(H_2|D)} = \frac{P(H_1)}{P(H_2)} \frac{P(D|H_1)}{P(D|H_2)}$$

Odds Ratios

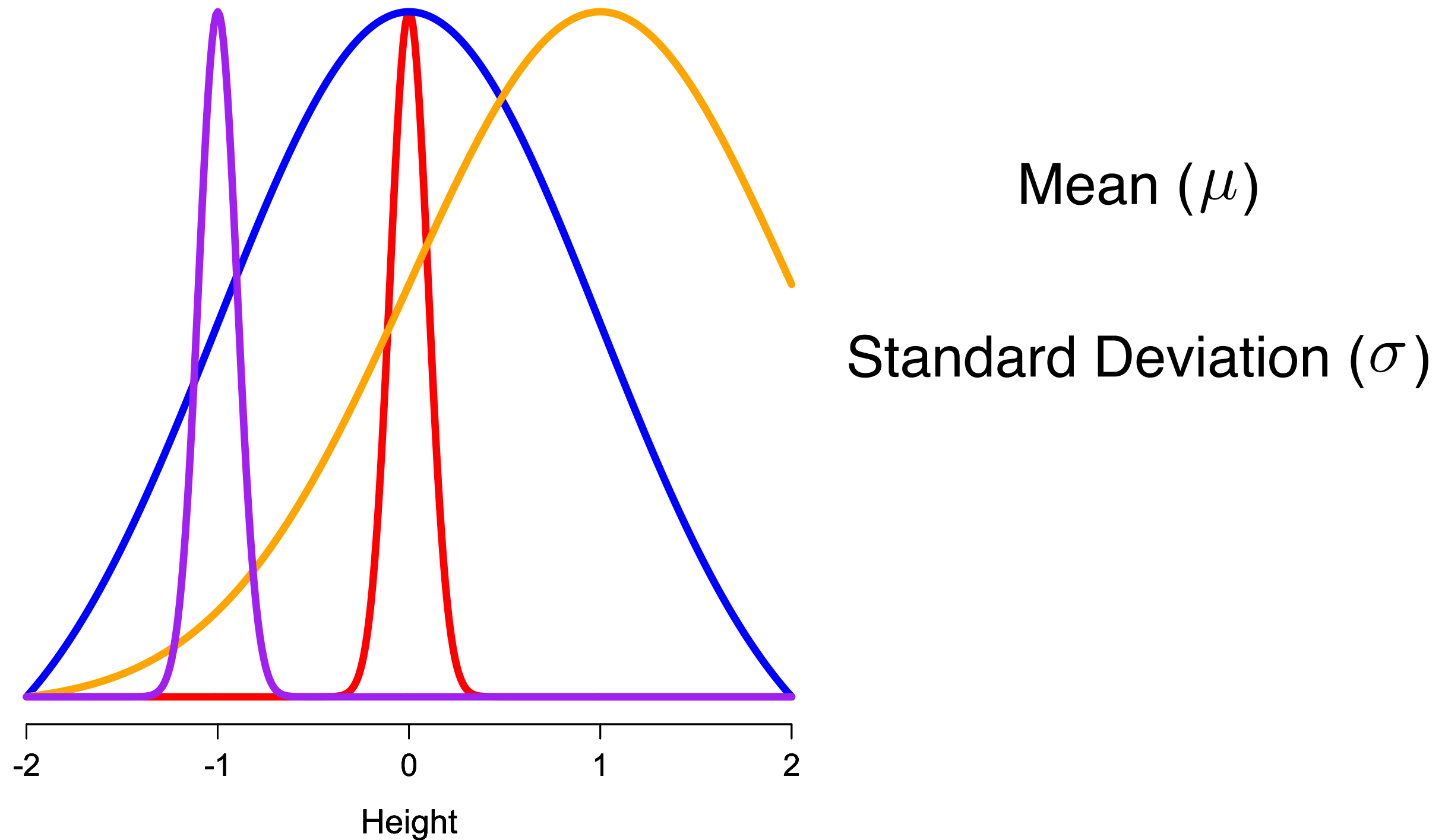
Prior Odds

Bayes Factor

Posterior Odds

$$\frac{P(H_1)}{P(H_2)} \times \frac{P(D|H_1)}{P(D|H_2)} = \frac{P(H_1|D)}{P(H_2|D)}$$

Estimating Parameters of a Normal Distribution



Marginalizing

Joint Probabilities

| | σ_1 | σ_2 | σ_3 | |
|---------|------------|------------|------------|------|
| μ_1 | 0.10 | 0.07 | 0.12 | 0.29 |
| μ_2 | 0.05 | 0.22 | 0.06 | 0.33 |
| μ_3 | 0.05 | 0.19 | 0.14 | 0.38 |
| | 0.20 | 0.48 | 0.32 | |

Marginal Probabilities

The diagram shows a 3x3 matrix of joint probabilities for variables μ_1, μ_2, μ_3 and $\sigma_1, \sigma_2, \sigma_3$. The joint probabilities are highlighted in blue. The marginal probabilities for each μ_i (sum of columns) and for each σ_j (sum of rows) are highlighted in red. An arrow points from the label 'Joint Probabilities' to the blue area, and two arrows point from the label 'Marginal Probabilities' to the red areas.

Monte Carlo Methods

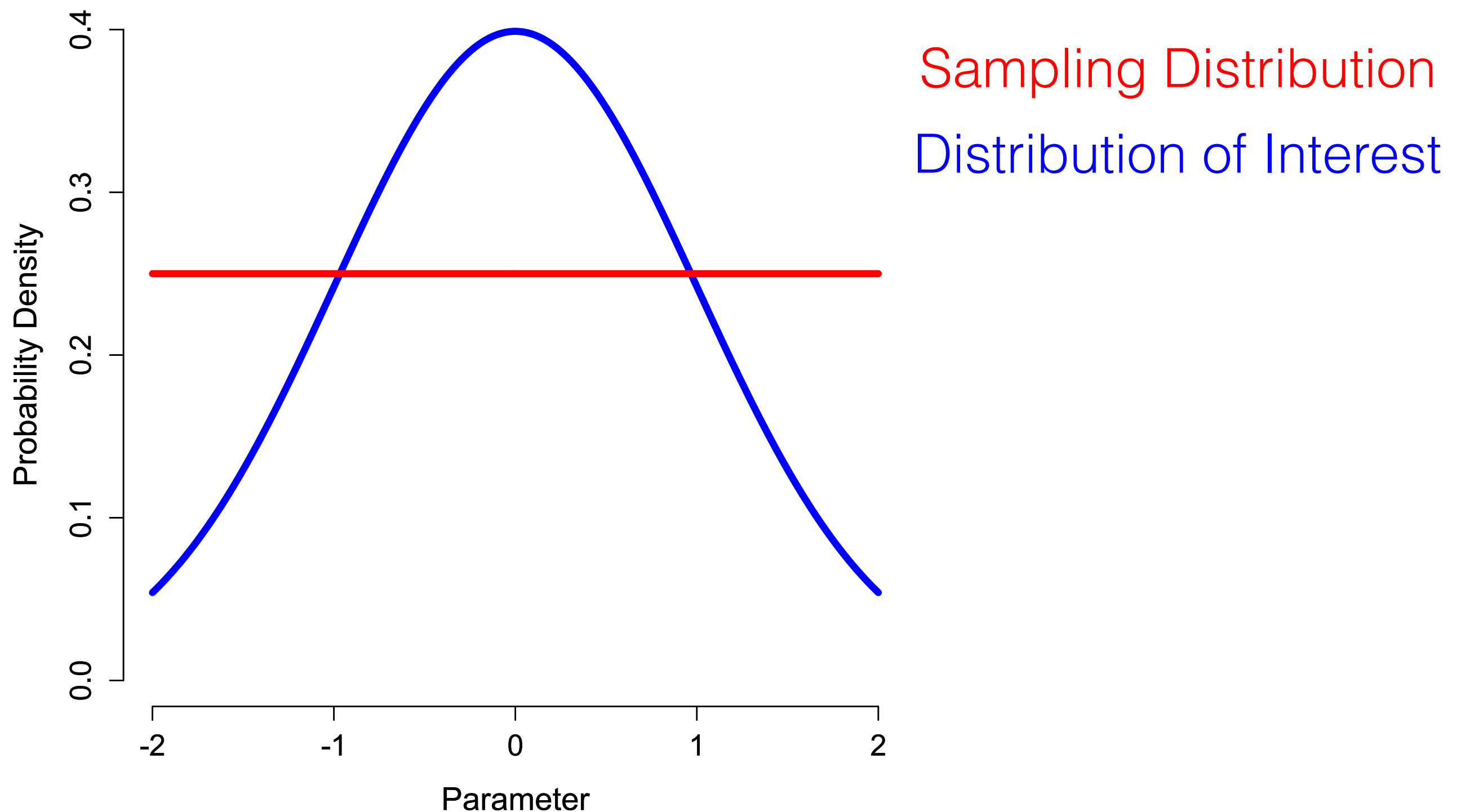
An entire class of methods to draw samples from or estimate moments (mean, variance, etc.) from distributions when closed-form solutions are not available. Rely on drawing (pseudo-)random numbers.



(Monaco, not Las Vegas)

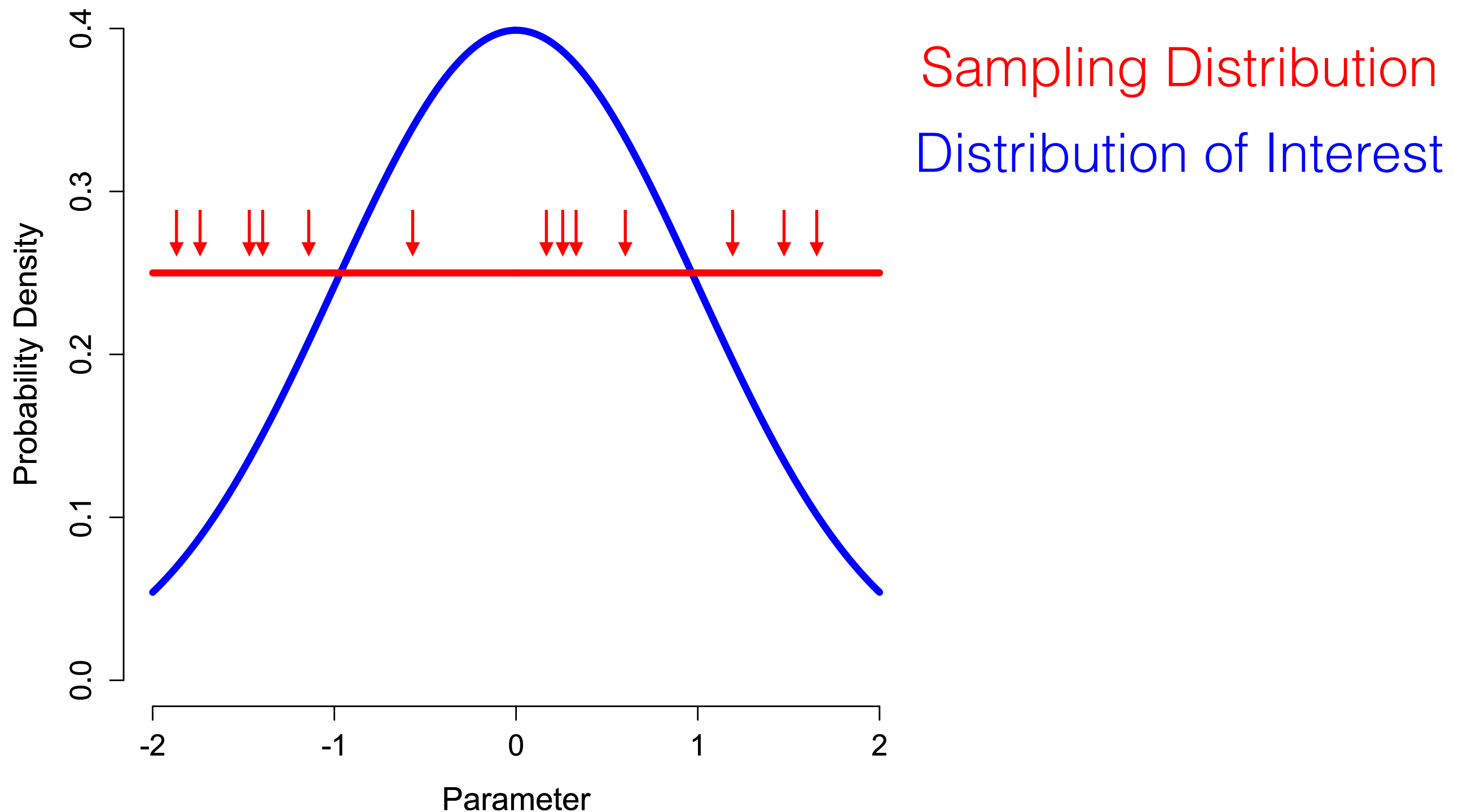
Monte Carlo Methods

Importance Sampling



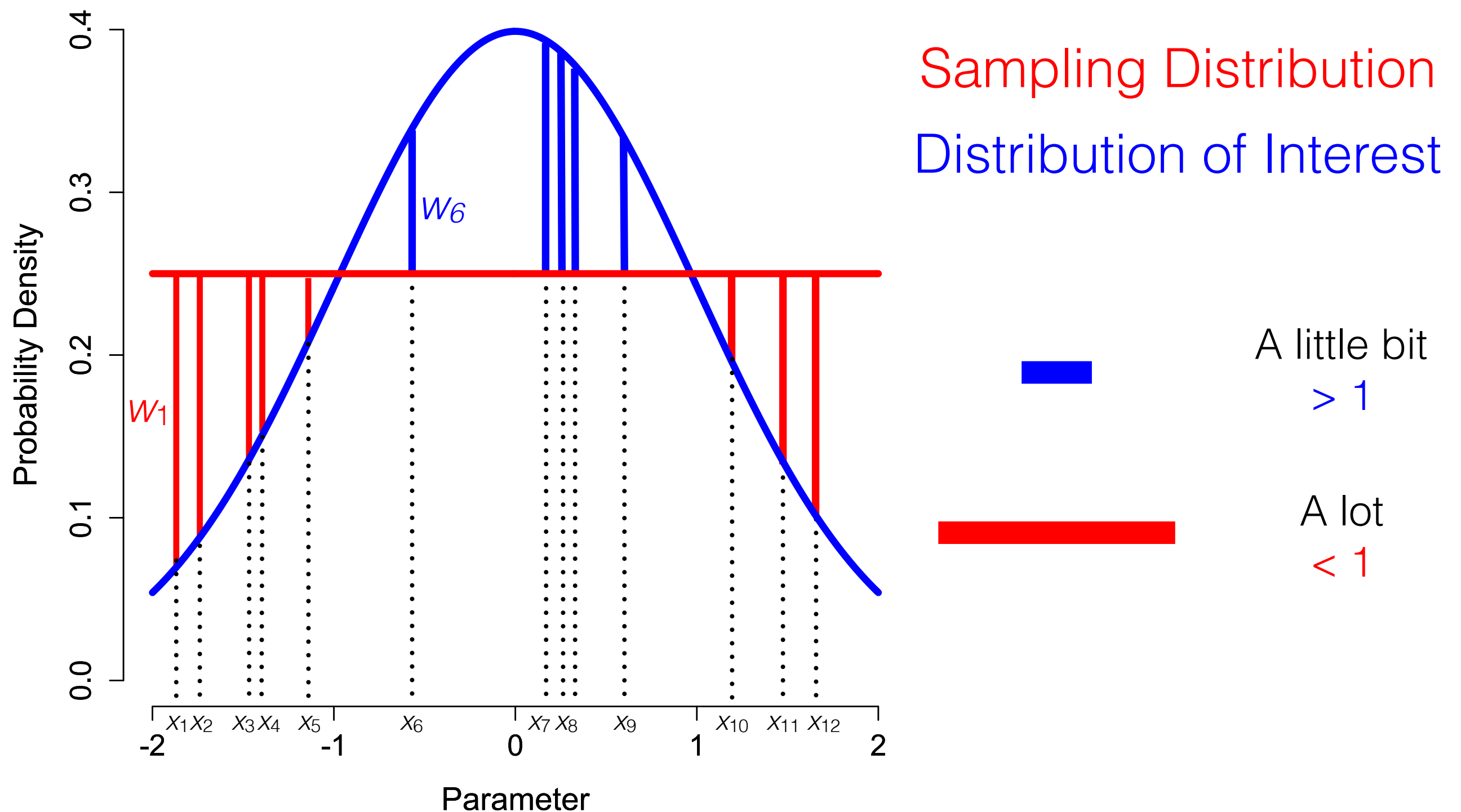
Monte Carlo Methods

Importance Sampling



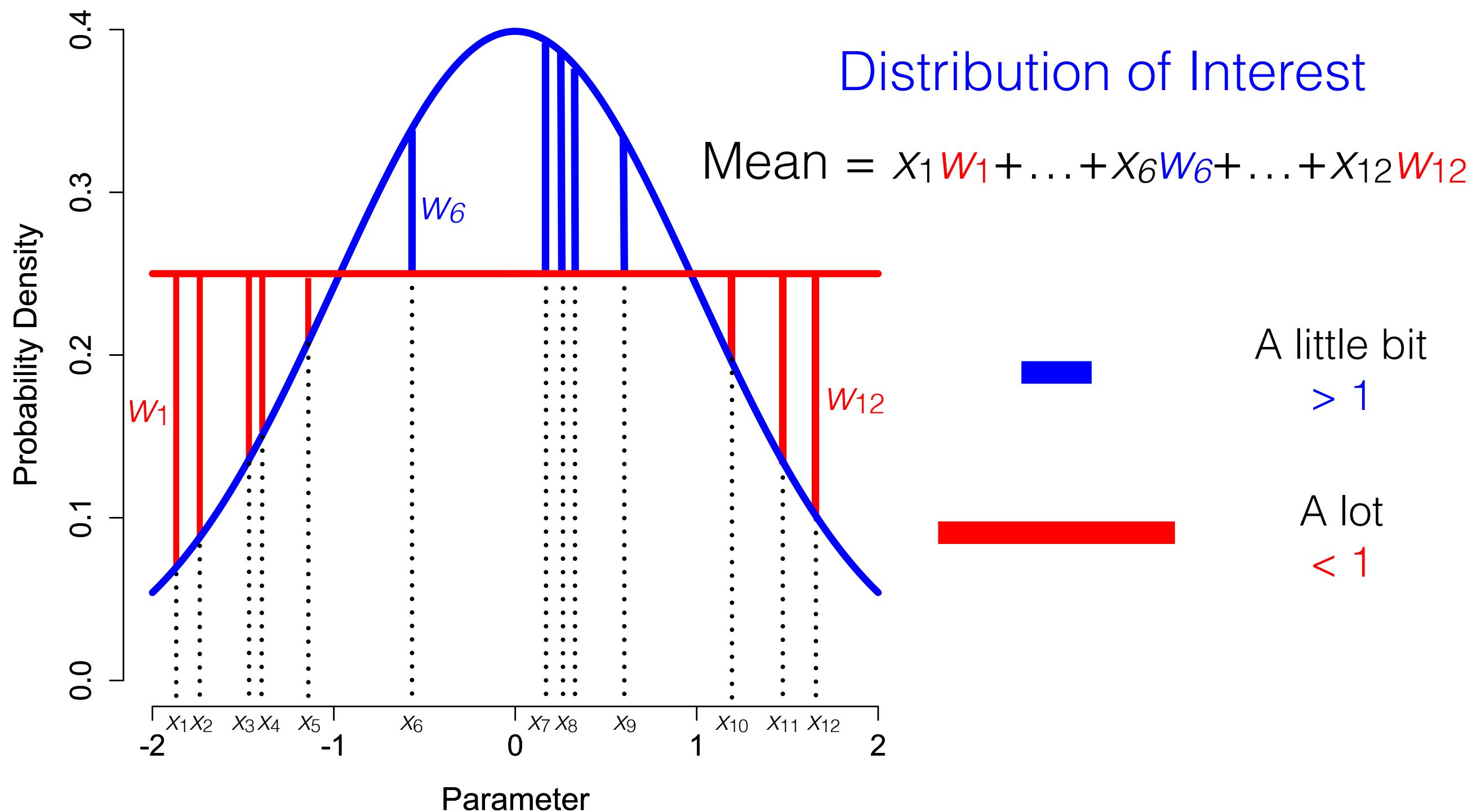
Monte Carlo Methods

Importance Sampling



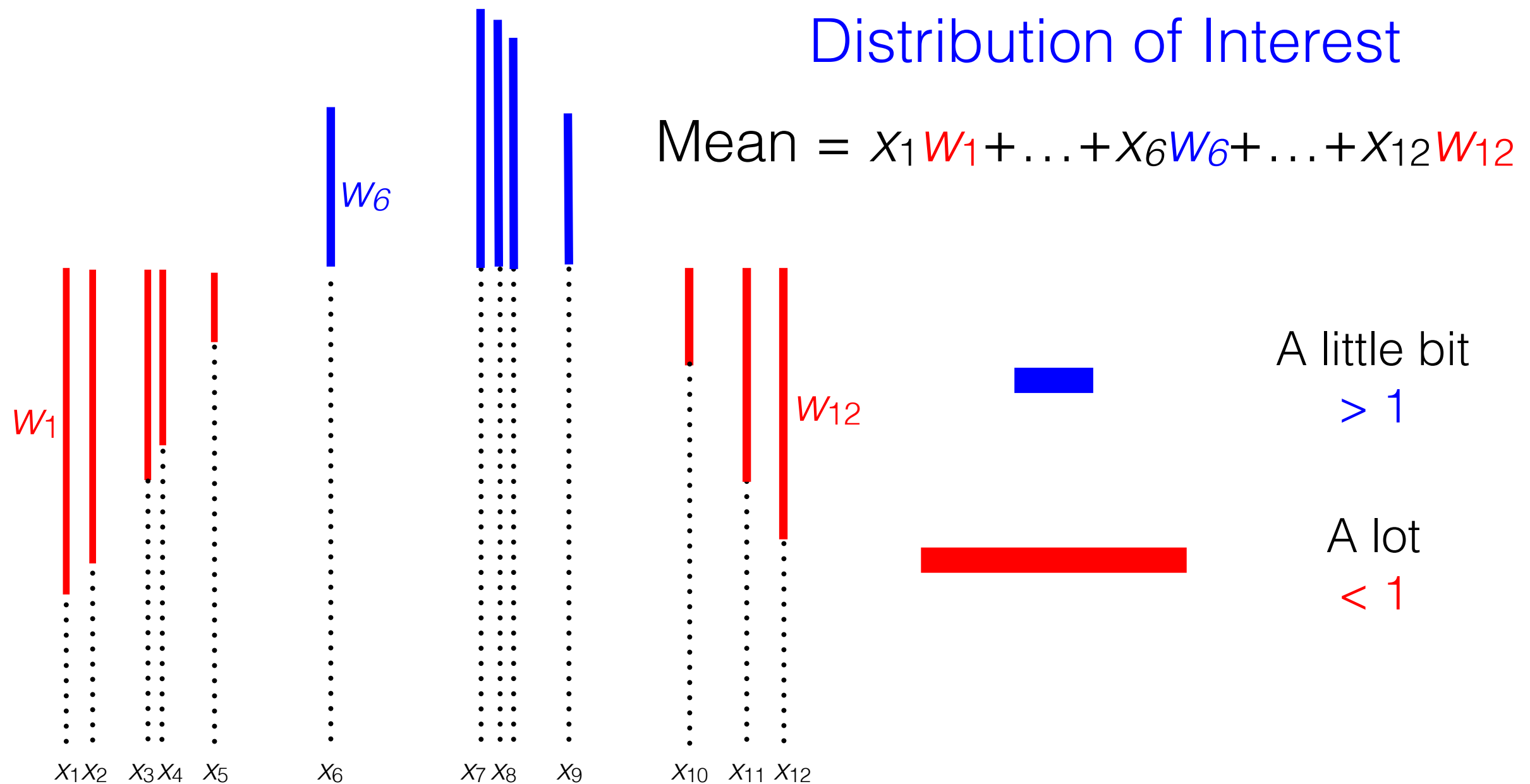
Monte Carlo Methods

Importance Sampling



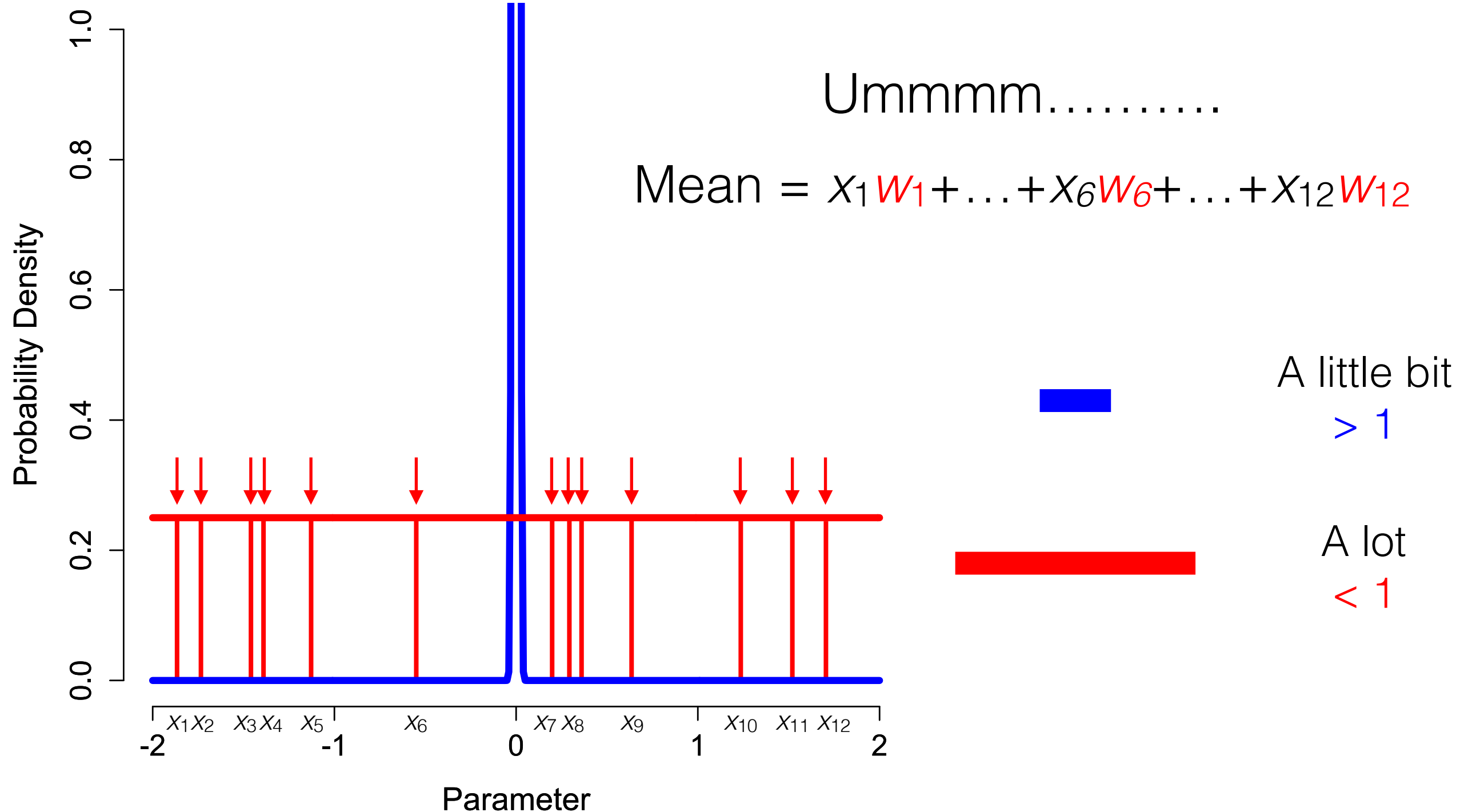
Monte Carlo Methods

Importance Sampling



Monte Carlo Methods

Importance Sampling

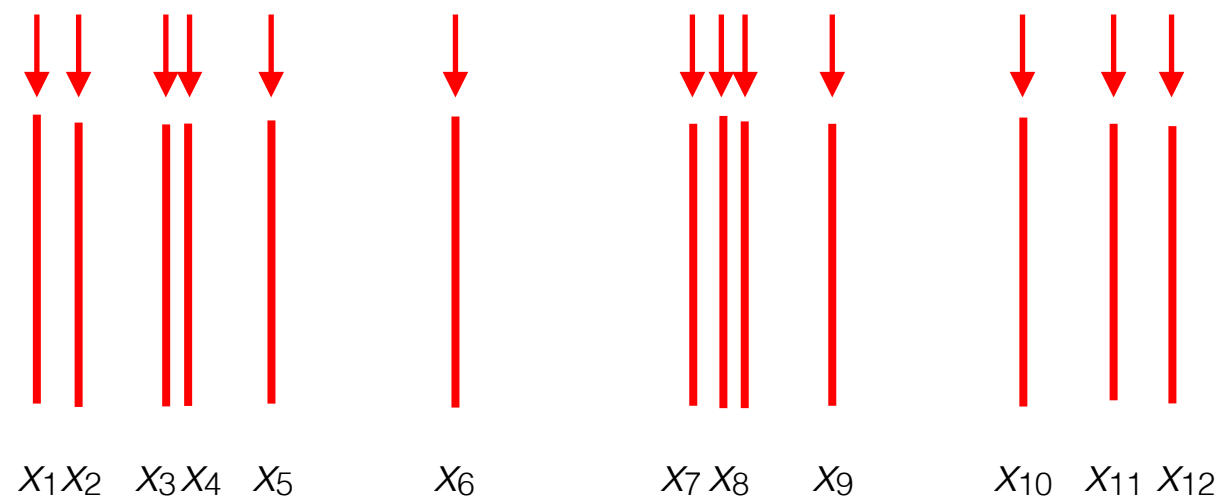


Monte Carlo Methods

Importance Sampling

Does this look like our
Distribution of Interest?

$$\text{Mean} = X_1 W_1 + \dots + X_6 W_6 + \dots + X_{12} W_{12}$$



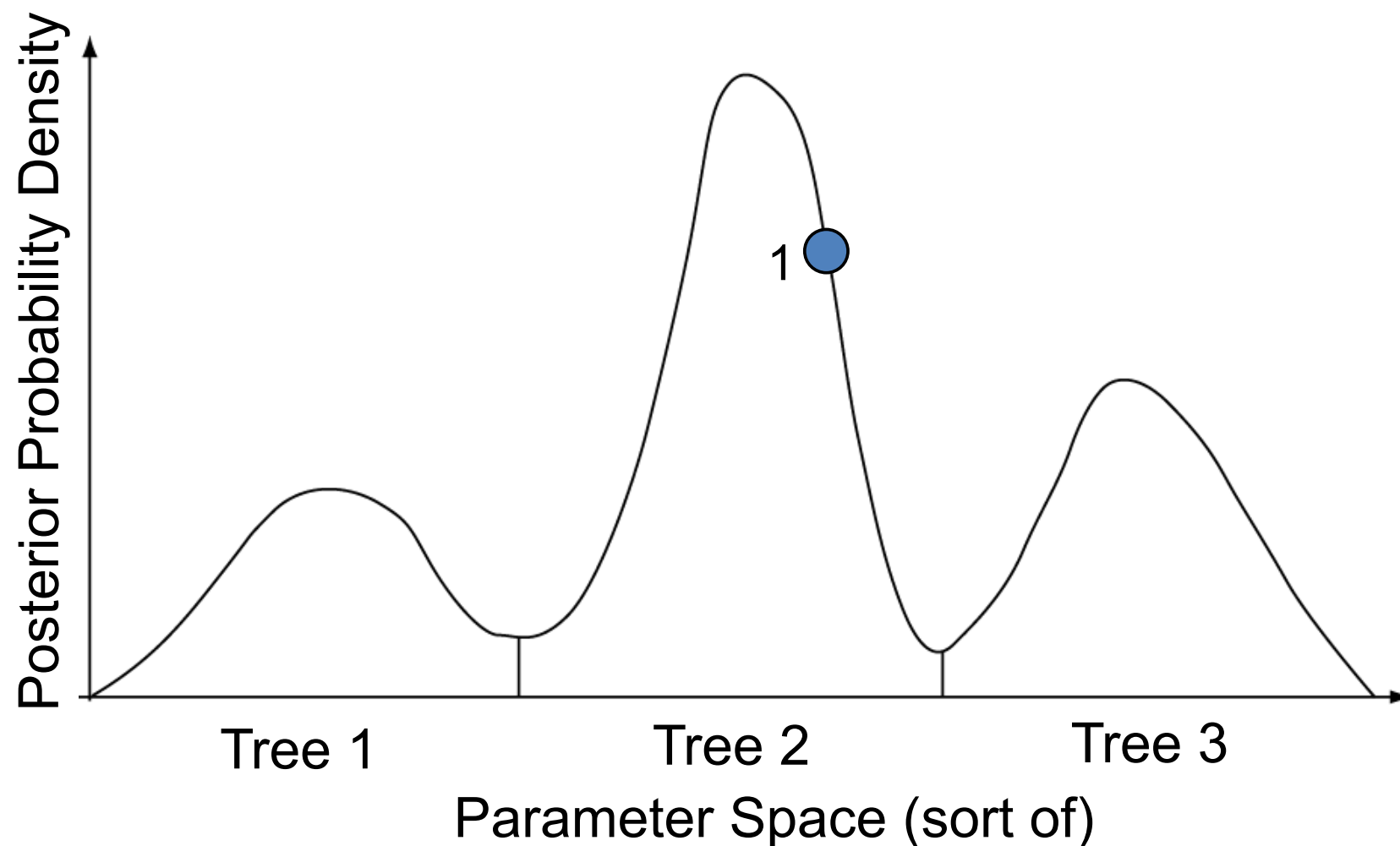
A little bit
 > 1



A lot
 < 1

Markov chain Monte Carlo

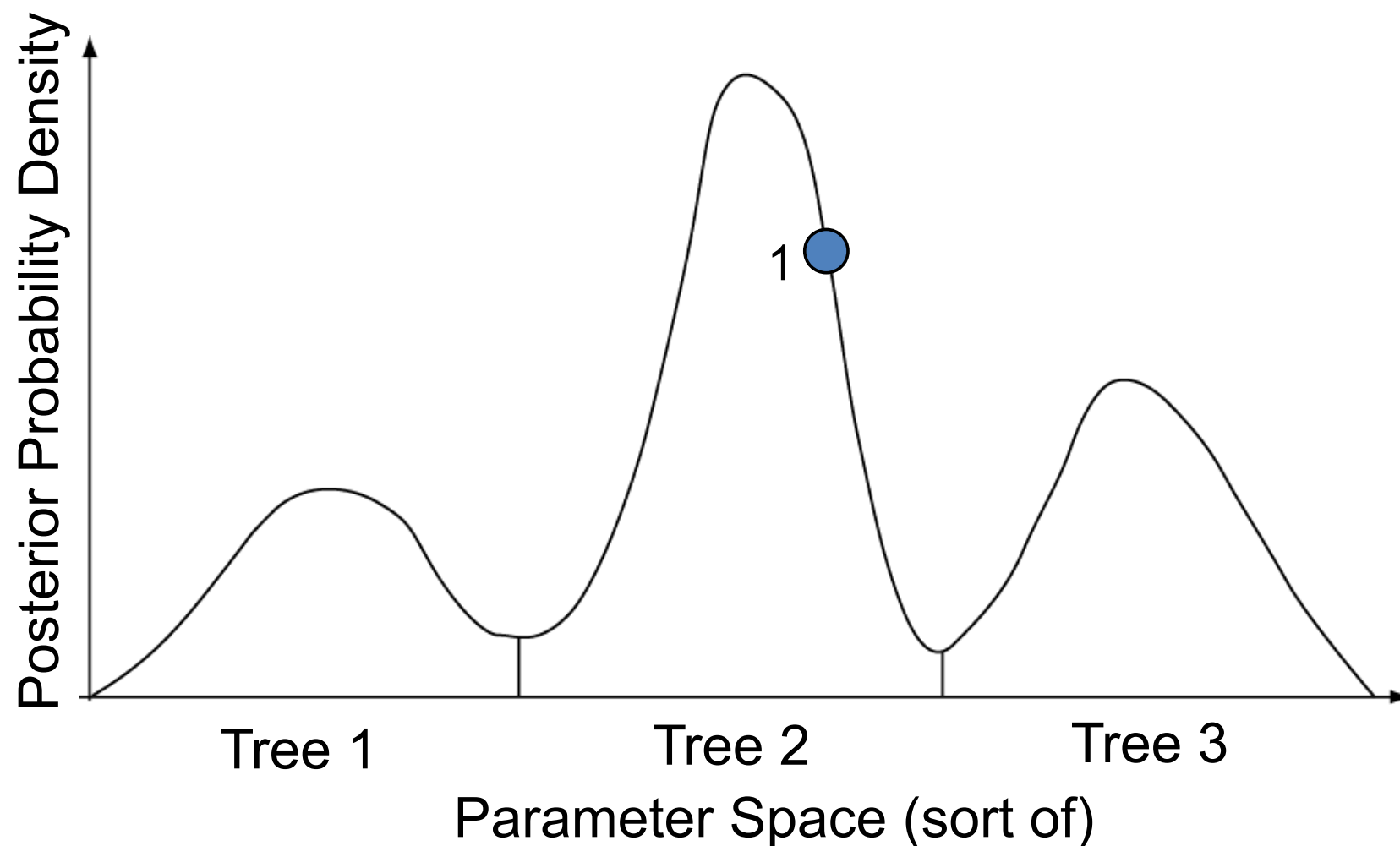
1. Start at an arbitrary point



This slide “borrowed” from F. Ronquist

Markov chain Monte Carlo

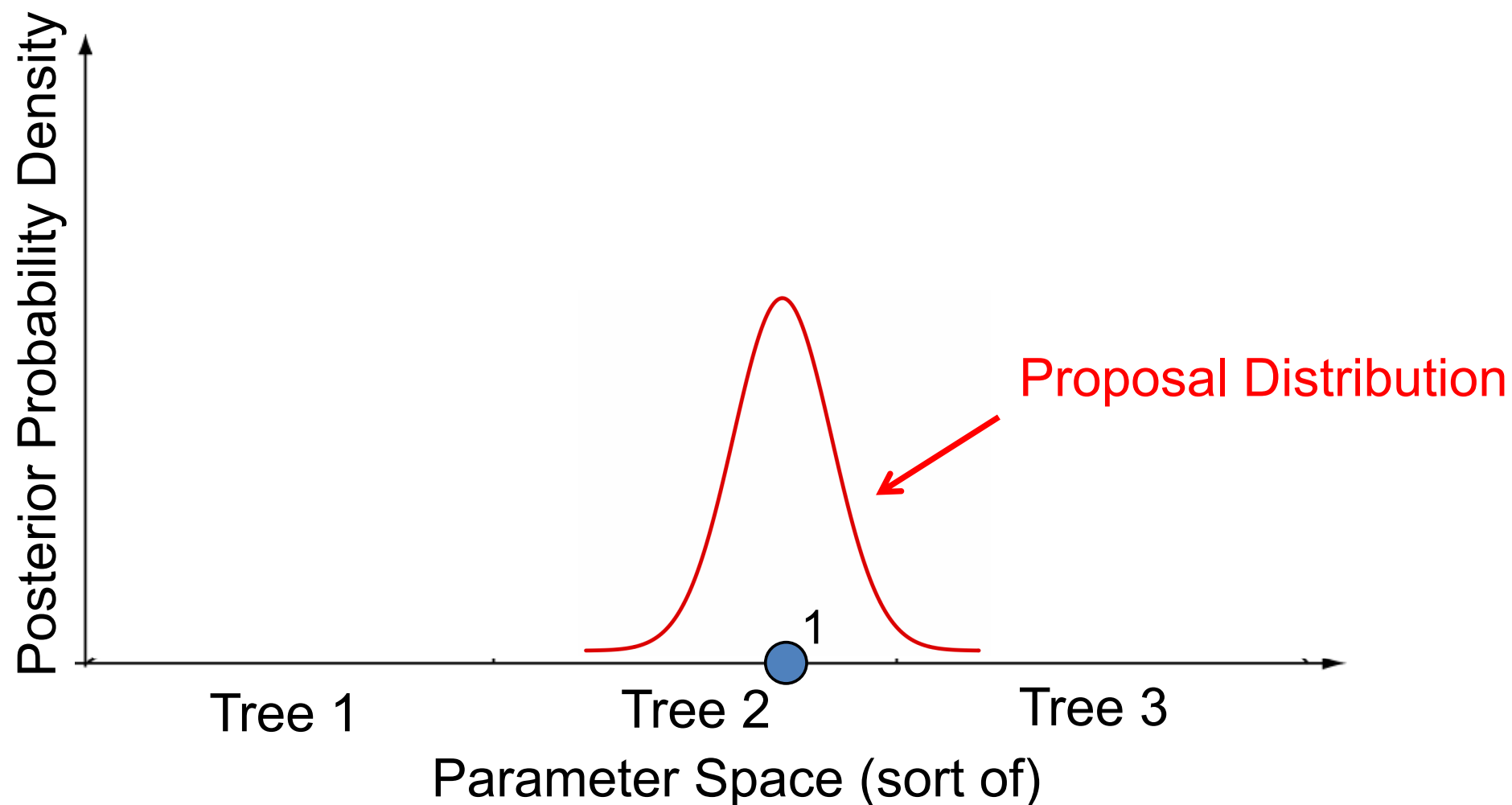
1. Start at an arbitrary point
2. Make a small random move



This slide “borrowed” from F. Ronquist

Markov chain Monte Carlo

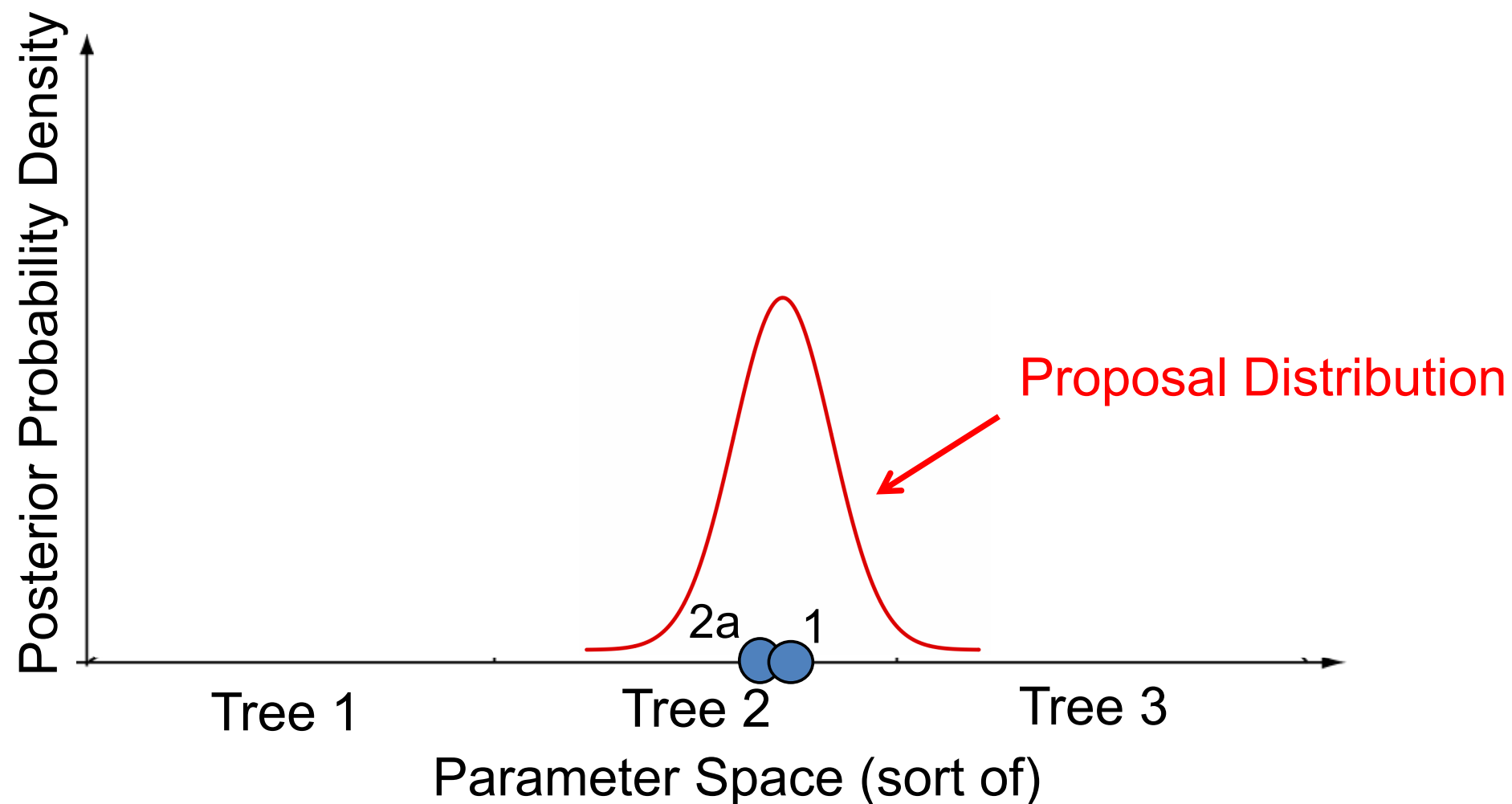
1. Start at an arbitrary point
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Markov chain Monte Carlo

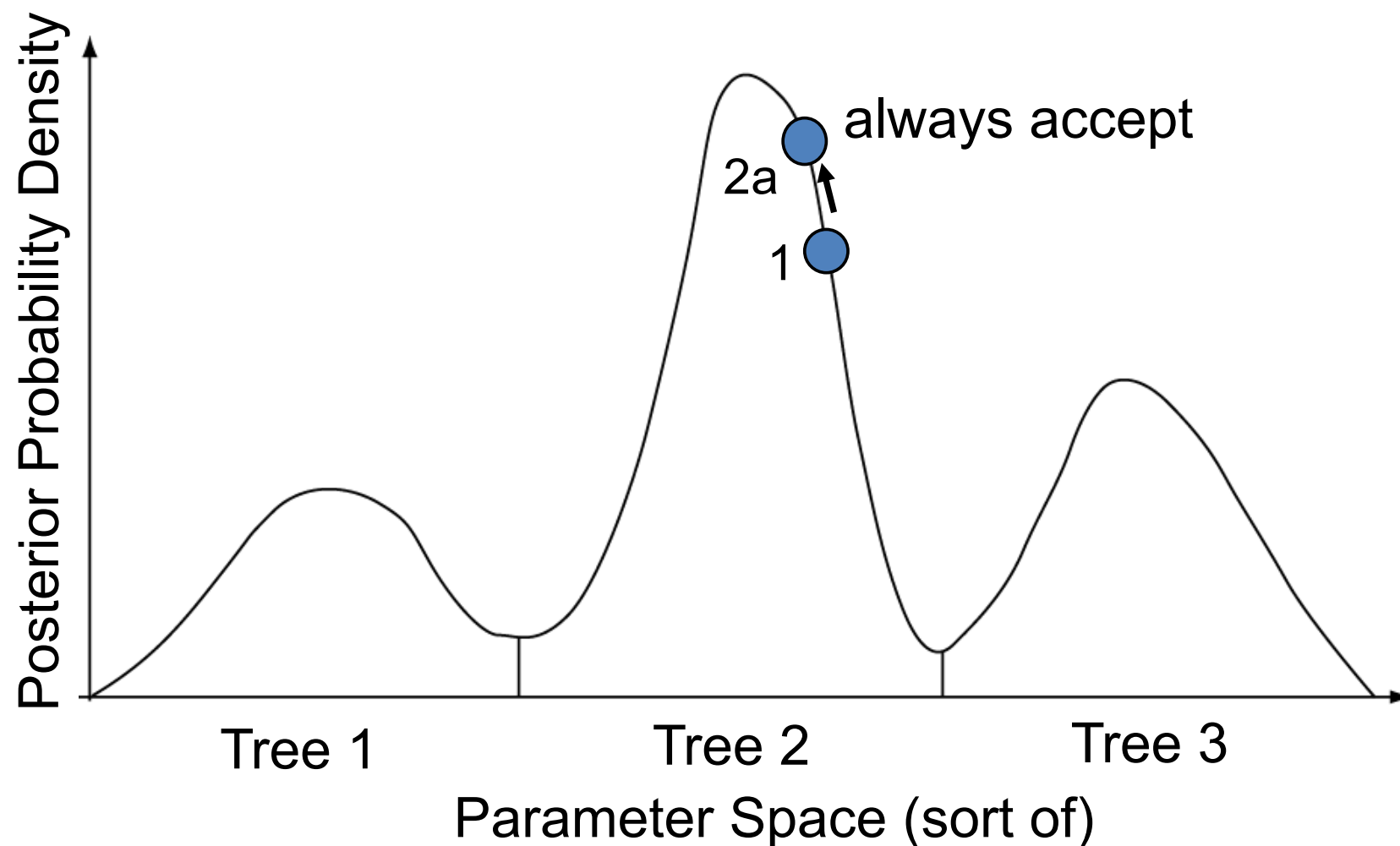
1. Start at an arbitrary point
2. Make a small random move



This slide “borrowed” from F. Ronquist

Markov chain Monte Carlo

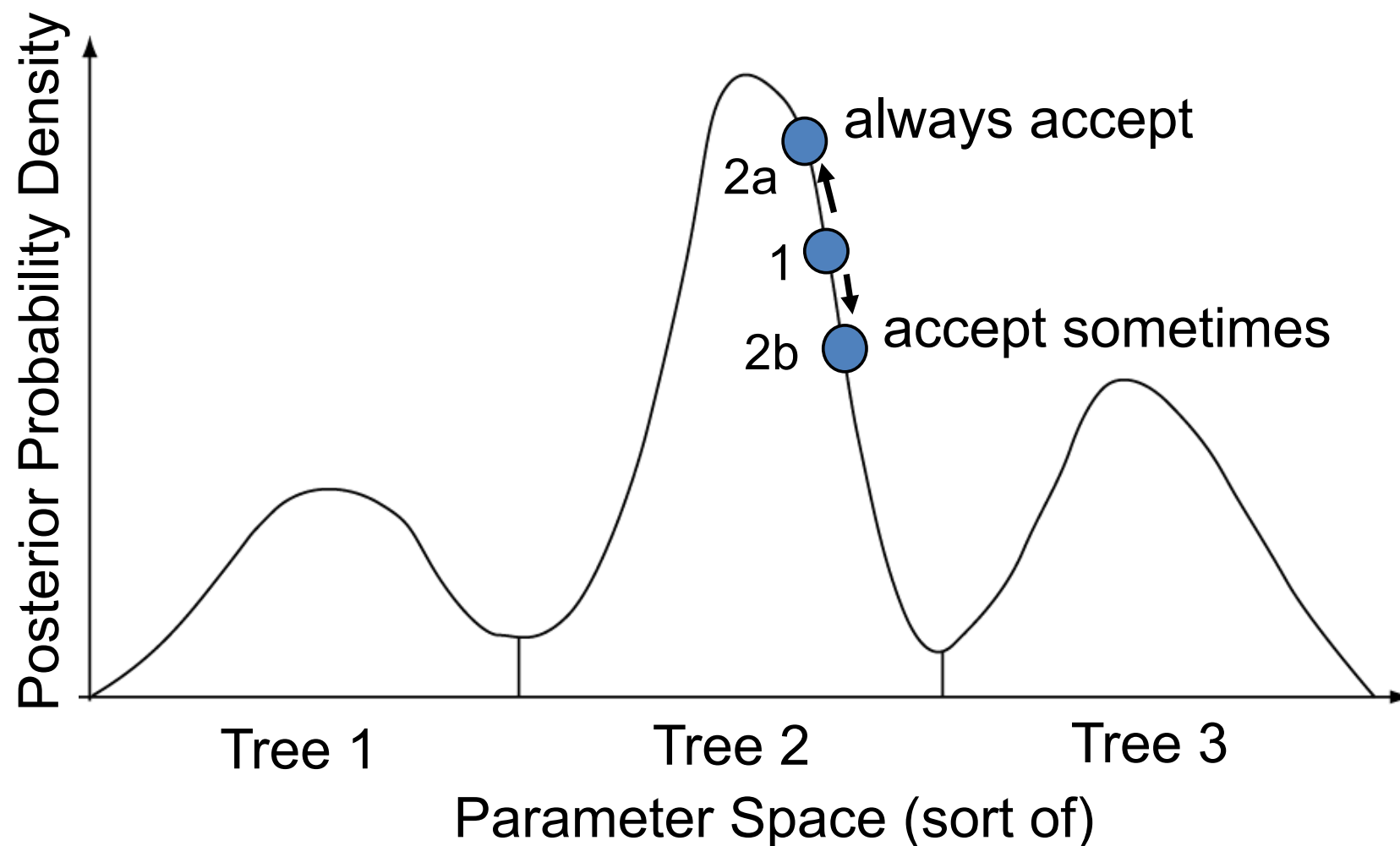
1. Start at an arbitrary point
2. Make a small random move
3. Calculate posterior density ratio (r) of new state to old state:
 - a) $r > 1 \rightarrow$ new state accepted
 - b) $r < 1 \rightarrow$ new state accepted with probability r . If new state not accepted, stay in the old state



This slide “borrowed” from F. Ronquist

Markov chain Monte Carlo

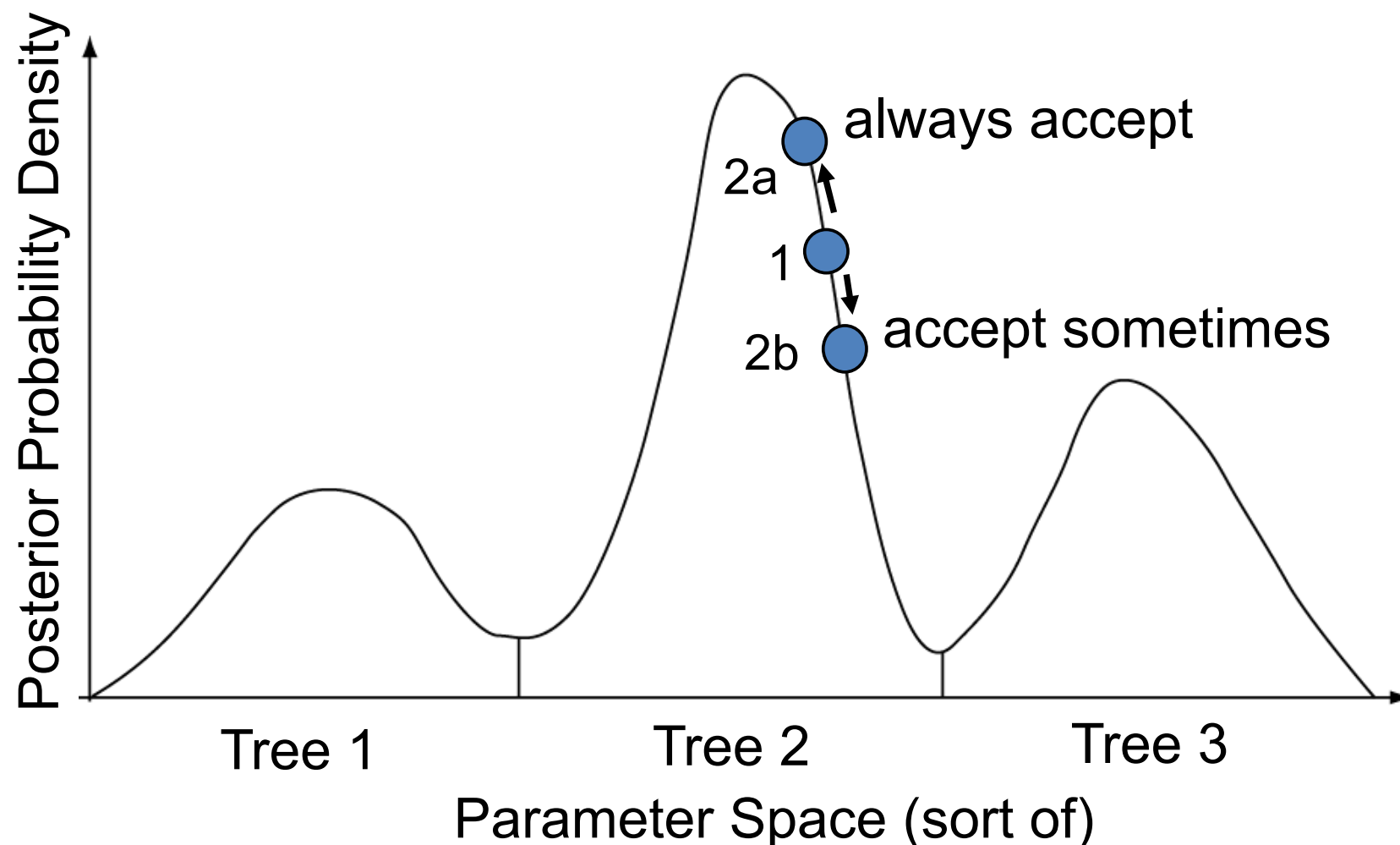
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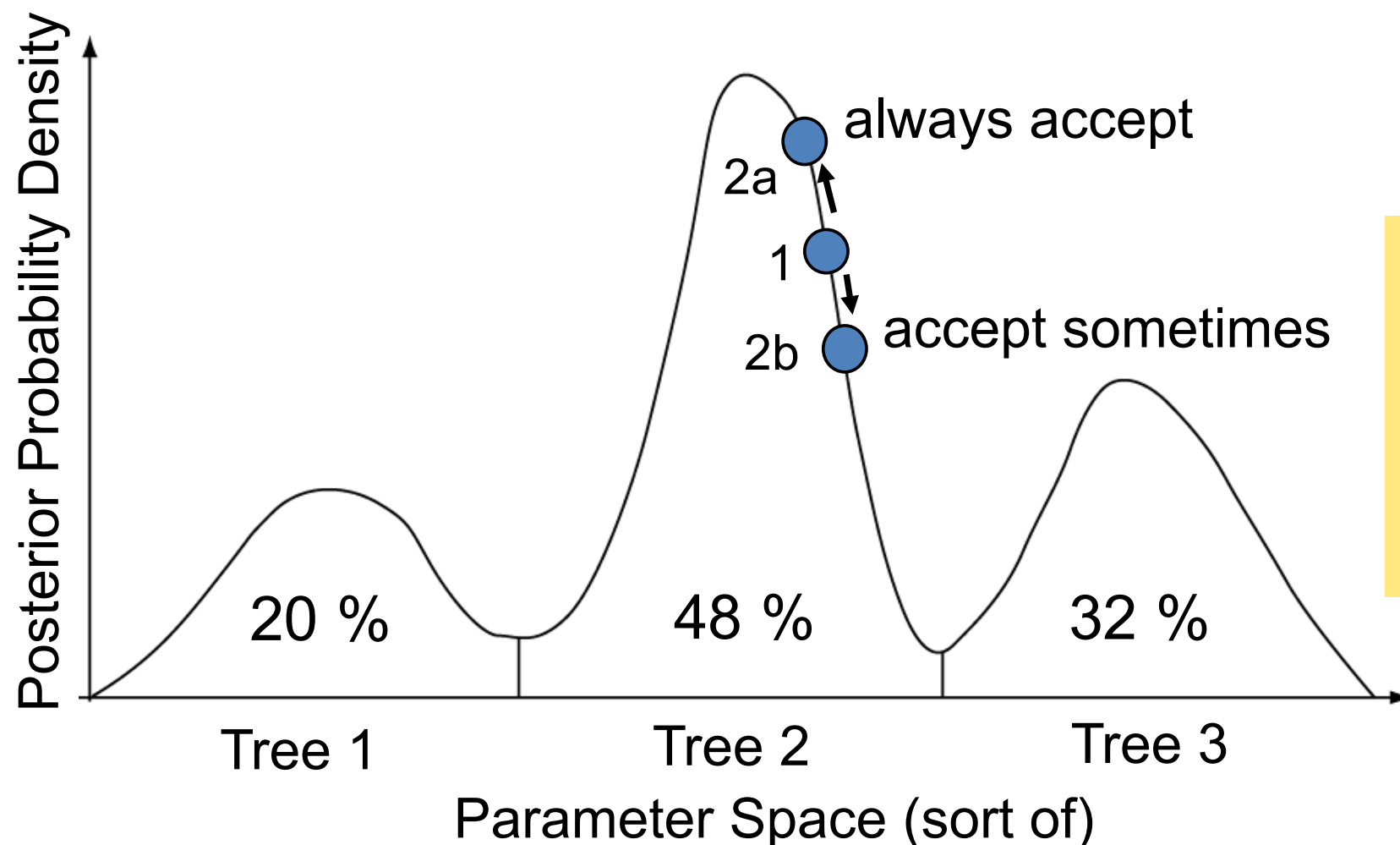
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4. Go to step 2 a BUNCH (x 10,000' s – x 10,000,000' s)



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Markov chain Monte Carlo

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The proportion of time the MCMC procedure samples from a particular parameter region is an estimate of that region's posterior probability density

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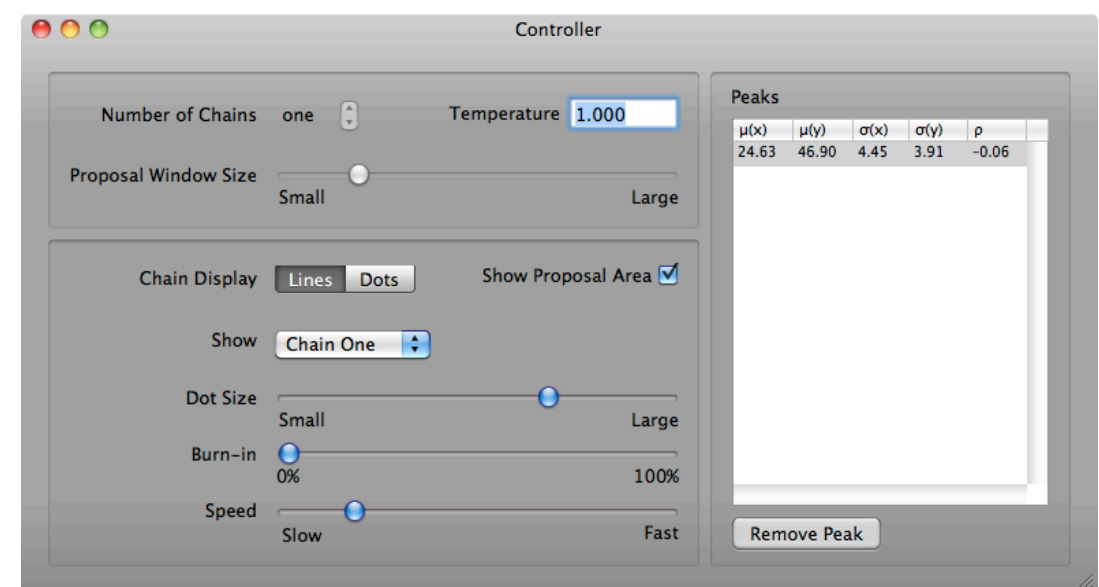
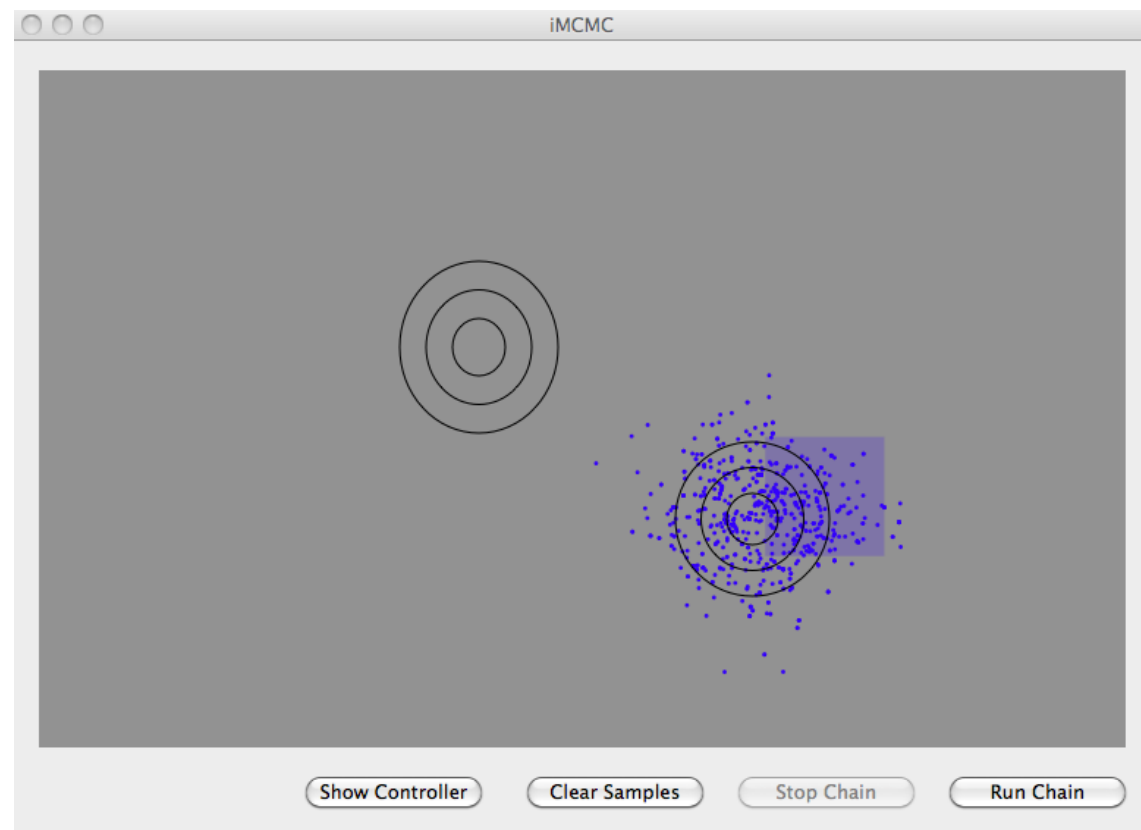
Toy MCMC Software

MCMC Robot (Paul Lewis; PC & iOS)

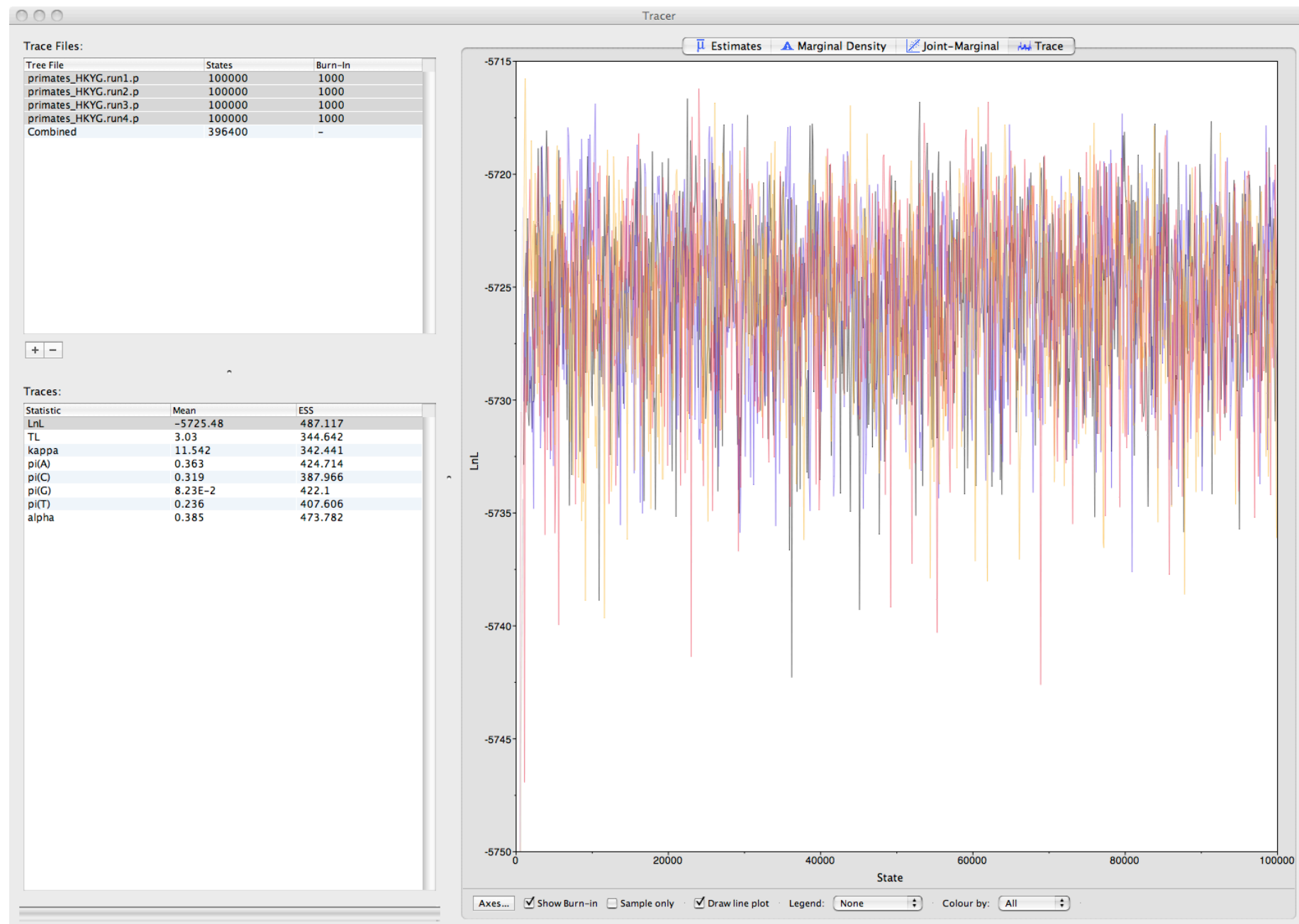
<http://www.mcmicrobot.org>

iMCMC (John Huelsenbeck; Mac)

<http://cteg.berkeley.edu/software/huelsenbeck/McmcApp.zip>



Convergence of Scalars - Tracer



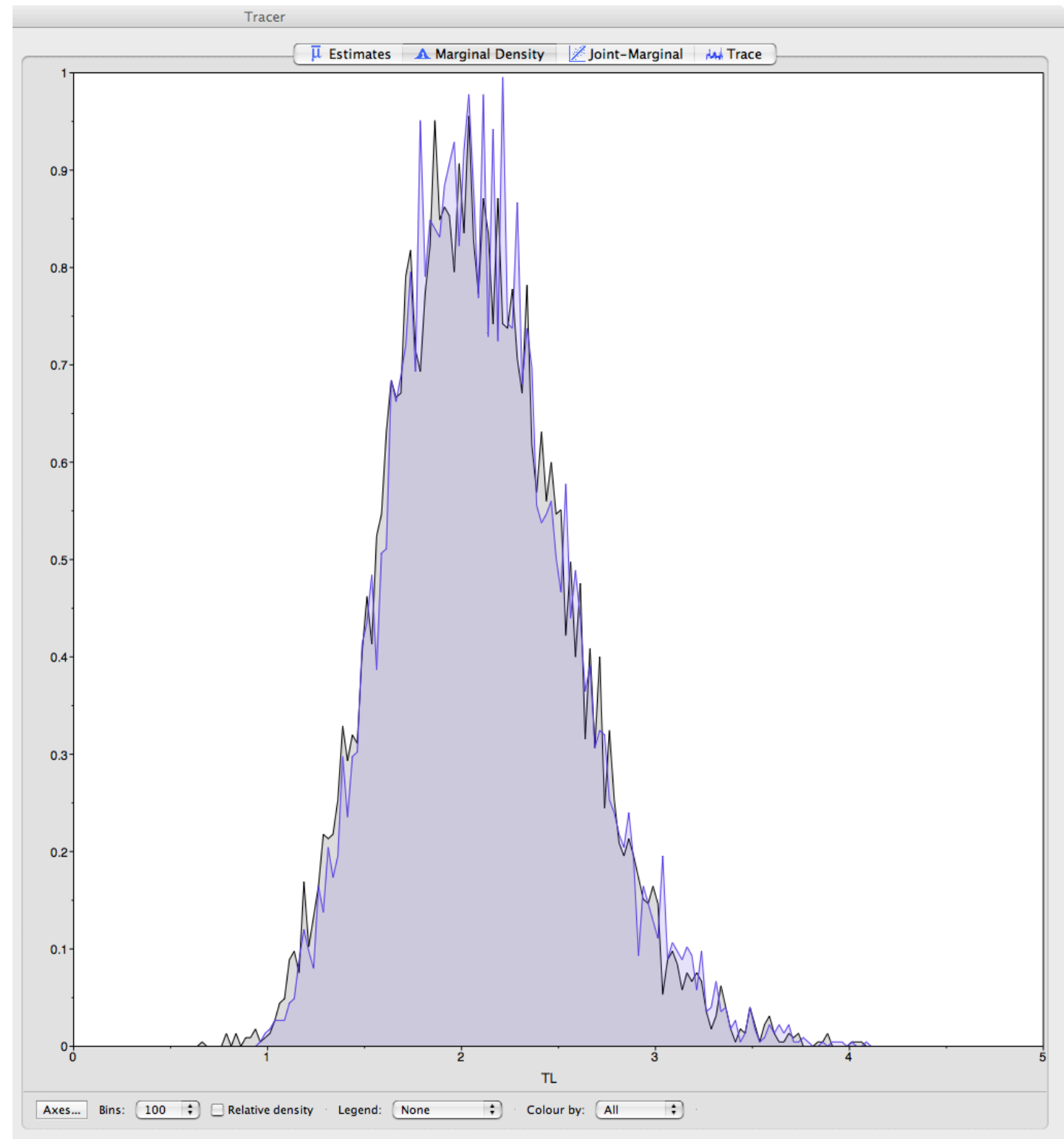
Running on Empty



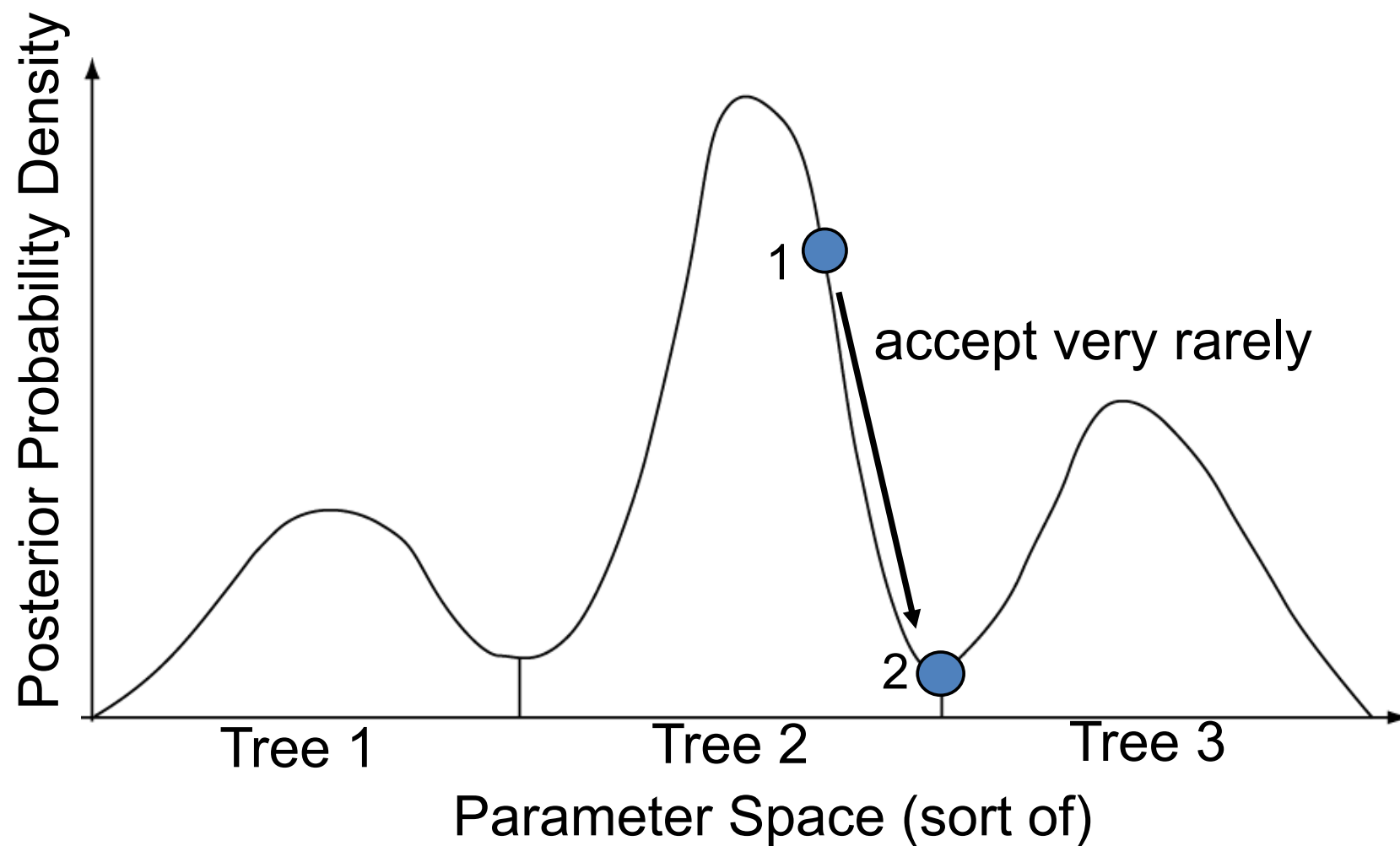
```
#NEXUS
begin data;
dimensions ntax=12 nchar=5;
format datatype=dna interleave=no gap=- missing=?;
matrix
Tarsius_syrichta      ??????
Lemur_catta           ??????
Homo_sapiens          ??????
Pan                   ??????
Gorilla               ??????
Pongo                 ??????
Hylobates             ??????
Macaca_fuscata        ??????
M_mulatta             ??????
M_fascicularis        ??????
M_sylvanus            ??????
Saimiri_sciureus      ??????
;
end;
```

Or now in MrBayes:

mcmc data=no



Metropolis Coupling

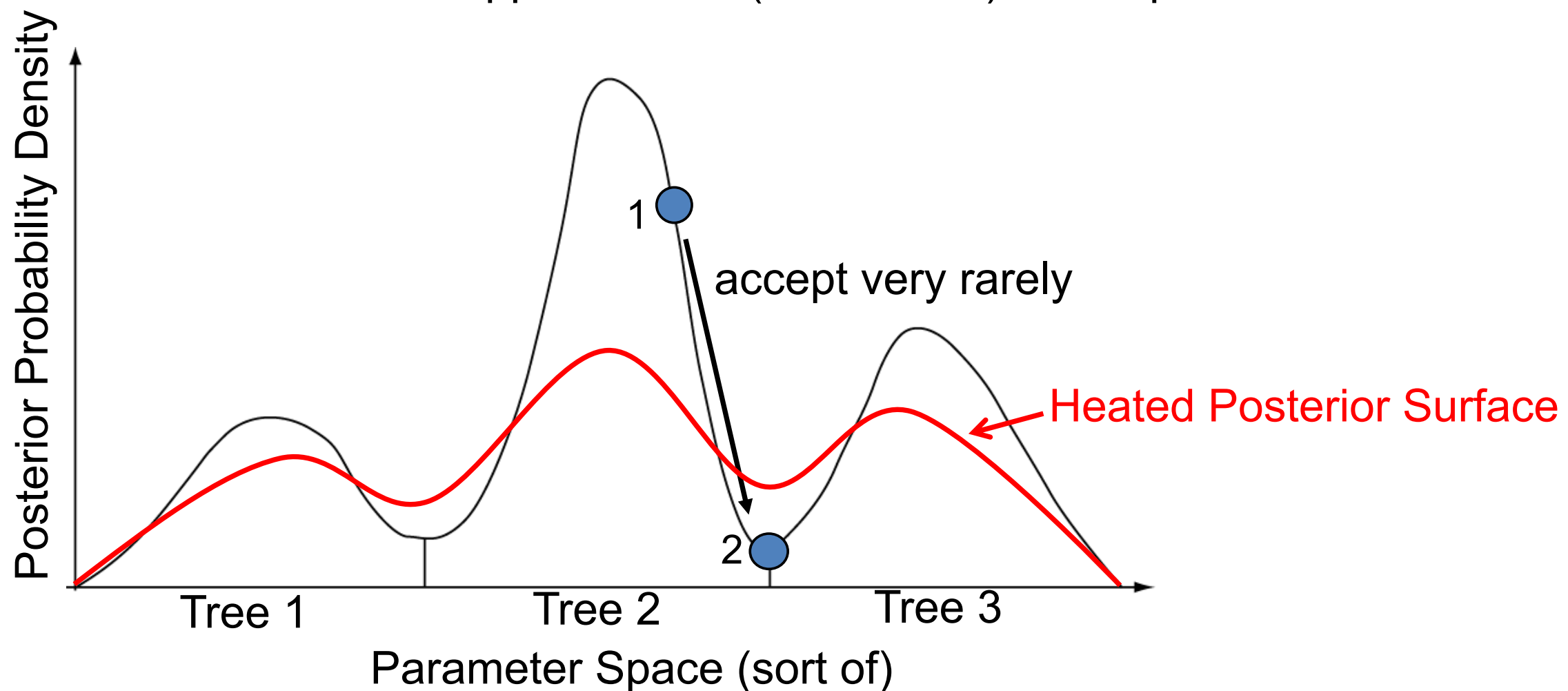


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Metropolis Coupling

- Same rules as regular MCMC, but now there are multiple chains with different ‘temperatures’.
- ‘Heated’ chains sample a ‘melted’ version of the posterior
- Only difference is that heated chains raise the ratio of posterior densities to $(1 - \text{temp})$ when deciding whether to accept a move.

$r^{(1-\text{temp})}$ approaches 1 (flat surface) as temp. increases

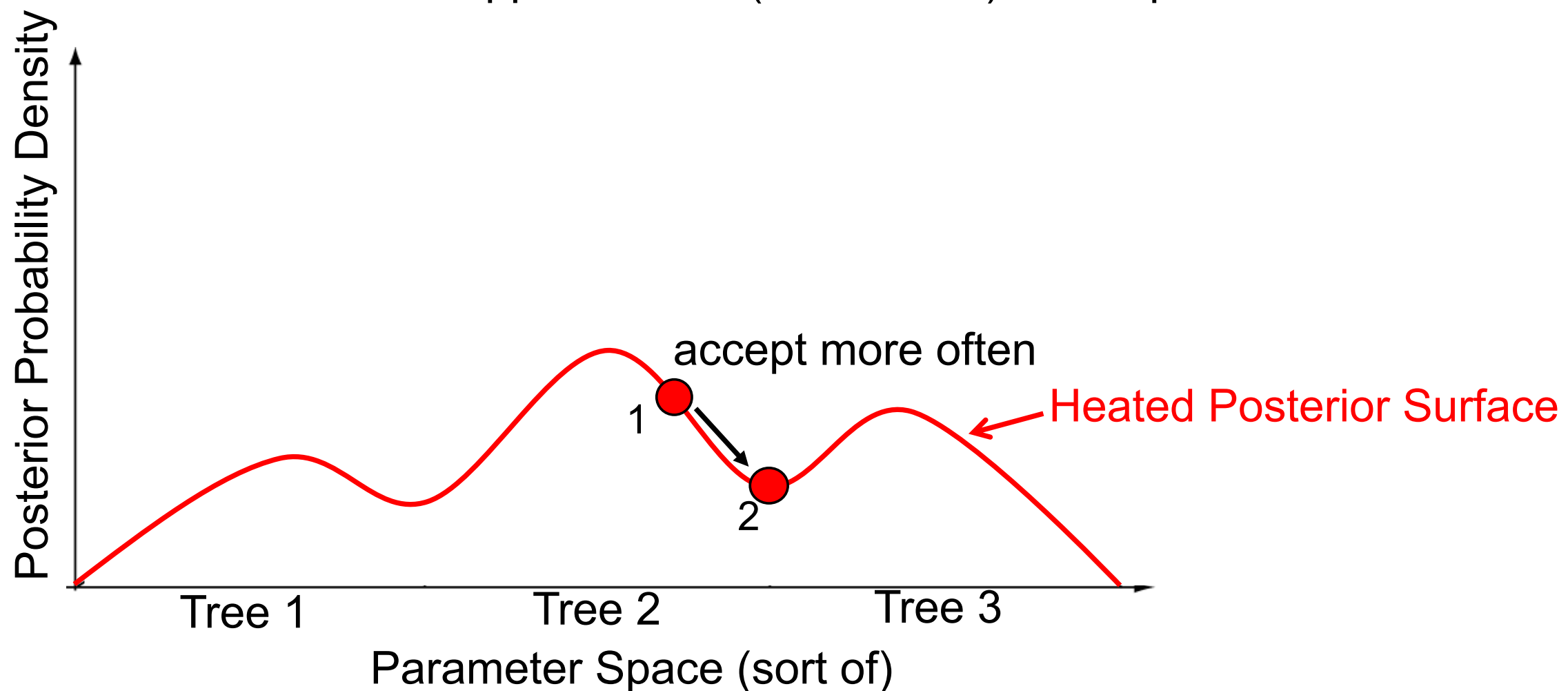


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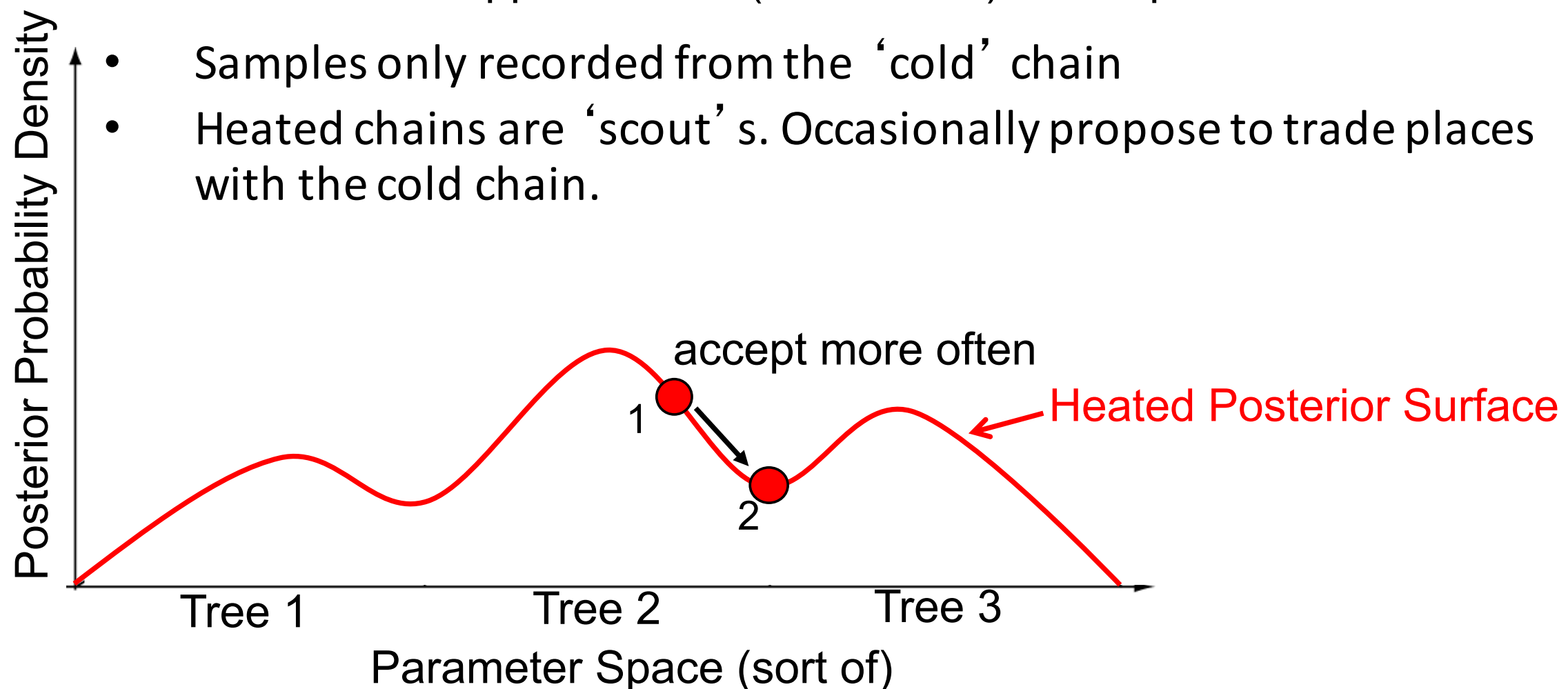
This slide “borrowed” from F. Ronquist

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$r^{(1-\text{temp})}$ approaches 1 (flat surface) as temp. increases

- Samples only recorded from the ‘cold’ chain
- Heated chains are ‘scout’ s. Occasionally propose to trade places with the cold chain.



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