



# Bayesian Sampling Methods for Population Genetics

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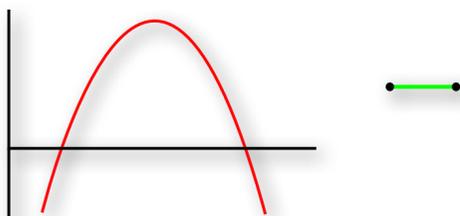


DEPARTMENT of SCIENTIFIC COMPUTING

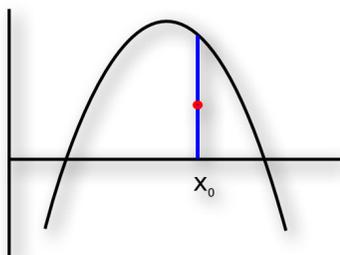
## Slice Sampling

as Done by Neil 2003

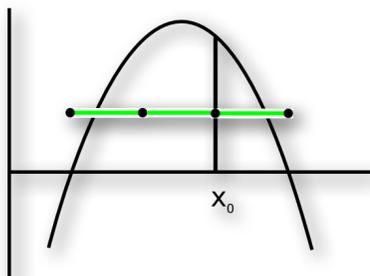
Start with a **function** to evaluate



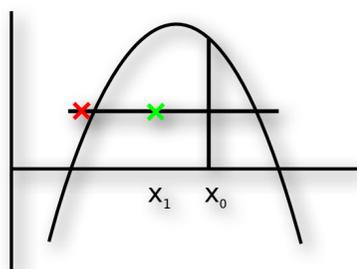
and a **window size**



Evaluate the function and choose a **random height**



Step out until outside of function is reached to create **slice**

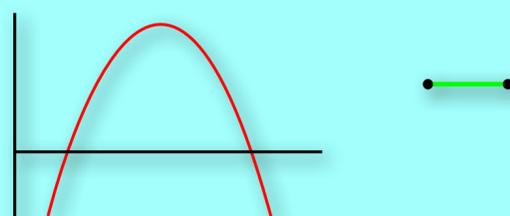


**New Point** is the first random uniform inside this "Slice"

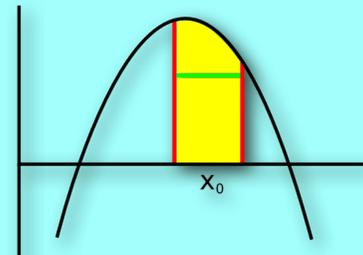
## Metropolis Hastings

sliding window proposal

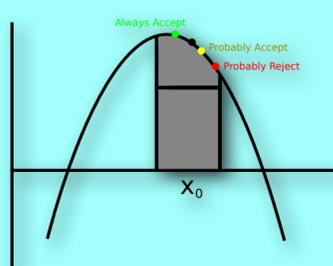
Start with a function to evaluate



and a **window size**



Choose a new point within **this window**

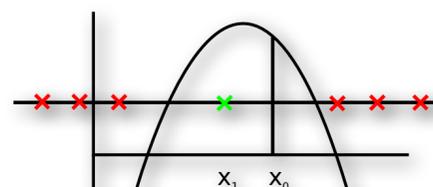


Evaluate this new point, and accept or reject it

$$\text{Probability of acceptance} = \min\left(1, \frac{P(X')}{P(X_0)}\right)$$

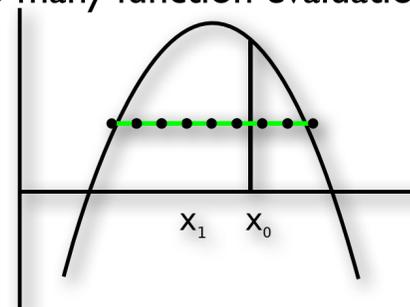
## Adaptive Slice Sampling

We noticed that the efficiency of this algorithm was dependent on window size



Too large window size leads to many rejections

Too big or too small a window leads to too many function evaluations



Too small a window size requires many steps to reach outside the function

Solution is to dynamically adjust window size.



Convergence is quick

1:41 Adaptive

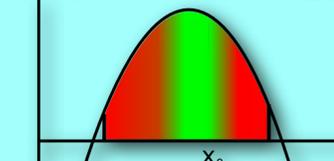


2:09 non-Adaptive

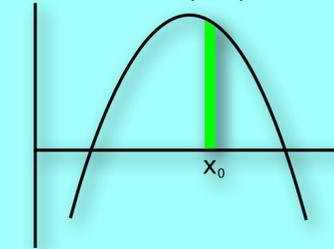
Result is quicker run time

## Adaptive Metropolis Hastings

The optimal acceptance rejection ratio is 0.44, depends on the window size



A large window leads to many samples that will be rejected



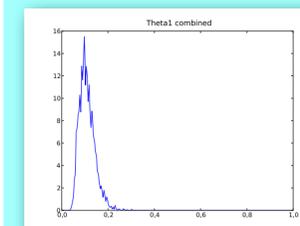
New samples aren't much different from old samples

too small or too large a window results in poor sampling

$$\beta = \left(\frac{1}{\alpha}\right)^{-\frac{R-1}{R}}$$

Rejection  $\rightarrow$  multiply window by beta  
Acceptance  $\rightarrow$  multiply window by alpha  
R  $\rightarrow$  desired acceptance/rejection ratio

We can choose our acceptance rejection ratio (must do this during burn in)

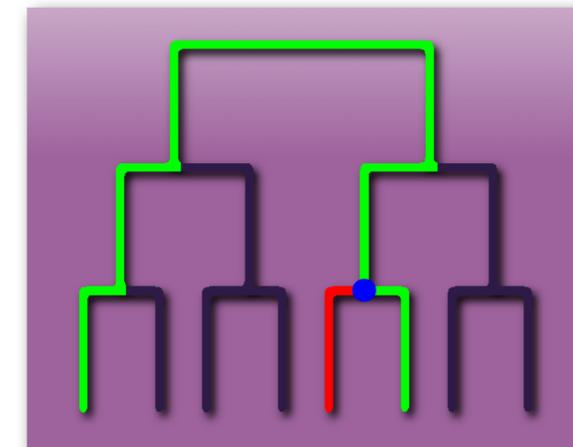


The adaptive algorithm(left) shows better mixing

Adaptive vs. Non adaptive

## Future Research

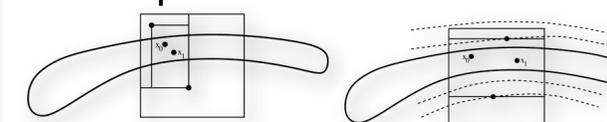
### Slice Sampling Trees



Slice sampling and adaptive MH on trees may be beneficial. Here I show one way to slice sample a tree. The red branch can be moved anywhere along the green line to reduce this type of sampling to one dimension at a time

## Multi-Dimensional

Currently sampling is done one parameter at a time. We know that these parameters are correlated



Multi dimensional slice sampling from Neil, 2003

$$Q_n(x, \cdot) = (1 - \beta)N(x, (2.38)^2 \Sigma_n / d) + \beta N(x, (0.1)^2 I_d / d)$$

Equation for adaptive multidimensional MH from Roberts and Rosenthal 2009

### References

Roberts GO and Rosenthal JS (2009) Examples of adaptive MCMC *Journal of Computational and Graphical Statistics* Vol 18(2) 349-367

Neil RM (2001) Slice Sampling *The Annals of Statistics* Vol 31(3) 705-767