METROPOLIS AND METROPOLIS-HASTINGS II

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ANNOUNCEMENTS

- Last homework (HW8) now online.
- Reminder: let the instructor know if you plan to request a letter grade.

OUTLINE

- Metropolis algorithm
 - Introduction and intuition
 - Algorithm
 - Illustration
 - Application to Poisson regression
- Metropolis-Hastings algorithm





INTRODUCTION

- As a refresher, suppose and suppose we specify a prior on .
- Then as usual, we are interested in

- As we already know, the challenge is that it is often difficult to compute
- Using the Monte Carlo method or Gibbs sampler, we have seen that we don't need to know. As long as we have conjugate and semiconjugate priors, we can generate samples directly from.
- What happens if we cannot sample directly from ?

MOTIVATING EXAMPLE

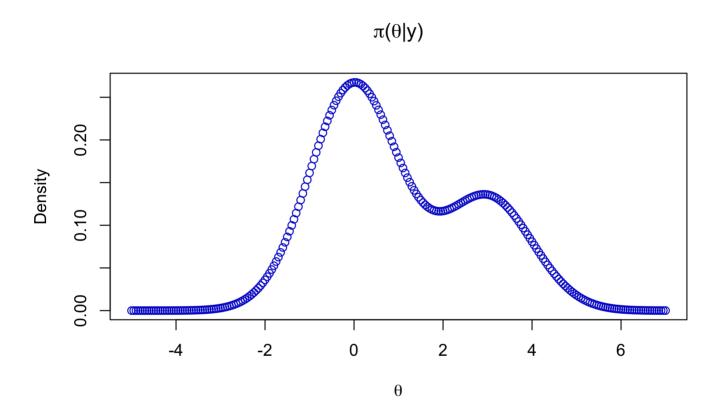
- To motivate our discussions on the Metropolis algorithm, let's explore a simple example.
- Suppose we wish to sample from the following density



- This is a mixture of two normal densities, one with mode near 0 and the other with mode near 3.
 - FYI: finite mixture models remains the most likely topic we will cover on Friday plus next part of next Wednesday.
- Anyway, let's use this density to explore the main ideas behind the Metropolis sampler.
- By the way, as you will see, we don't actually need to know the normalizing constant for Metropolis sampling but for this example, find it for practice! I will set it up in class.

MOTIVATING EXAMPLE

Let's take a look at the (normalized) density:



■ There are other ways of sampling from this density, but let's focus specifically on the Metropolis algorithm here.



- From a sampling perspective, we need to have a large group of values,
 from whose empirical distribution approximates
- That means that for any any two values and , we want

- Basically, we want to make sure that if and are in , the ratio of the number of the values equal to them properly approximates ——.
- How might we construct a group like this?

- Suppose we have a working group at iteration , and need to add a new value .
- Consider a candidate value that is close to (we will get to how to generate the candidate value in a minute). Should we set or not?
- By the way, notice that



which does not depend on the marginal likelihood we don't know!



- If
 - Intuition: is already a part of the density we desire and the density at is even higher than the density at .
 - Action: set
- = If ,
 - Intuition: relative frequency of values on our group equal to should be ______. For every , include only a fraction of an instance of .
 - Action: set with probability and with probability .

- This is the basic intuition behind the Metropolis algorithm.
- Where should the proposed value come from?
- Sample close to the current value using a symmetric proposal distribution . is actually a "family of proposal distributions", indexed by the specific value of .
- Here, symmetric means that
- The symmetric proposal is usually very simple with density concentrated near , for example, or .
- After obtaining , either add it or add a copy of to our current set of values, depending on the value of .



and setting

	The algorithm proceeds as tollows:					
	1. Given	, generate a candidate value	•			
	2. Compute the acceptance ratio					
	3. Set					
	which can be a	ccomplished by sampling	independently			



- Once we obtain the samples, then we are back to using Monte Carlo approximations for quantities of interest.
- That is, we can again approximate posterior means, quantiles, and other quantities of interest using the empirical distribution of our sampled values.

Some notes:

- The Metropolis chain ALWAYS moves to the proposed at iteration if has higher target density than the current .
- Sometimes, it also moves to a value with lower density in proportion to the density value itself.
- This leads to a random, Markov process than naturally explores the space according to the probability defined by , and hence generates a sequence that, while dependent, eventually represents draws from .



METROPOLIS ALGORITHM: CONVERGENCE

- We will not cover the convergence theory behind Metropolis chains in detail, but below are a few notes for those interested:
 - The Markov process generated under this condition is ergodic and has a limiting distribution.
 - Here, think of ergodicity as meaning that the chain can move anywhere at each step, which is ensured, for example, if everywhere!
 - By construction, it turns out that the Metropolis chains are reversible,
 so that convergence to is assured.
 - Think of reversibility as being equivalent to symmetry of the joint density of two consecutive and in the stationary process, which we do have by using a symmetric proposal distribution.
- If you want to learn more about convergence of MCMC chains, consider taking one of the courses on stochastic processes, or Markov chain theory.



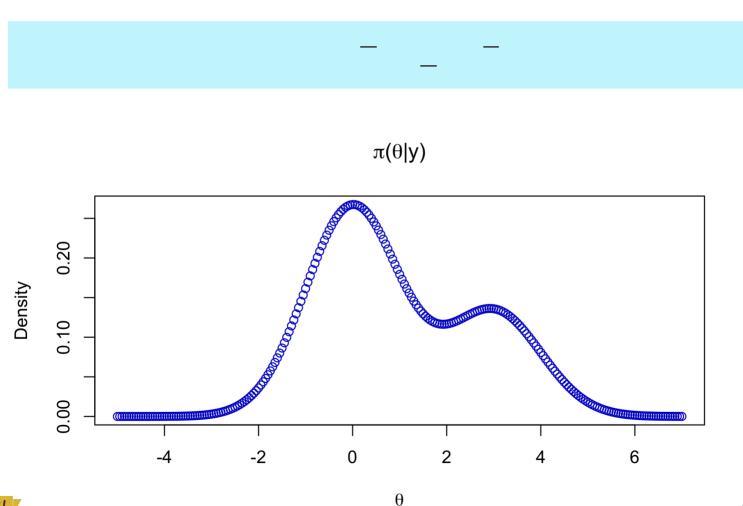
METROPOLIS ALGORITHM: TUNING

- Correlation between samples can be adjusted by selecting optimal (i.e., spread of the distribution) in the proposal distribution
- Decreasing correlation increases the effective sample size, increasing rate of convergence, and improving the Monte Carlo approximation to the posterior.
- However,
 - too small leads to for most proposed values, a high acceptance rate, but very small moves, leading to highly correlated chain.
 - too large can get "stuck" at the posterior mode(s) because can get very far away from the mode, leading to a very low acceptance rate and again high correlation in the Markov chain.
- Thus, good to implement several short runs of the algorithm varying and settle on one that yields acceptance rate in the range of 25-50%.
 - Burn-in (and thinning) is even more important here!



METROPOLIS IN ACTION

Back to our example with



IN-CLASS ANALYSIS: MOVE TO THE R SCRIPT HERE.



Poisson regression



COUNT DATA

- We will use the Metropolis sampler on count data with predictors, so let's first do some general review.
- Suppose you have count data as your response variable.
- For example, we may want to explain the number of c-sections carried out in hospitals using potential predictors such as hospital type, (that is, private vs public), location, size of the hospital, etc.
- The models we have covered so far are not (completely) adequate for count data with predictors.
- Of course there are instances where linear regression, with some transformations (especially taking logs) on the response variable, might still work reasonably well for count data.
- That's not the focus here, so we won't cover that.

Poisson regression

- As we have seen so far, a good distribution for modeling count data with no limit on the total number of counts is the Poisson distribution.
- As a reminder, the Poisson pmf is given by

Remember that

- When our data fails this assumption, we may have what is known as over-dispersion and may want to consider the Negative Binomial distribution instead (actually easy to fit within the Bayesian framework!).
- With predictors, index with , so that each is a function of .
 Therefore, the random component of the glm is

Poisson regression

 We must ensure that at any value of , therefore, we need a link function that enforces this. A natural choice is

- Combining these pieces give us our full mathematical representation for the Poisson regression.
- Clearly, has a natural interpretation as the "expected count", and

so the 's are multiplicative effects on the expected counts.

■ For the frequentist version, in R, use the glm command but set the option family = "poisson".

- We have data from a study of nesting horseshoe crabs (J. Brockmann, Ethology, 102: 1–21, 1996). The data has been discussed in Agresti (2002).
- Each female horseshoe crab in the study had a male crab attached to her in her nest.
- The study investigated factors that affect whether the female crab had any other males, called satellites, residing nearby her.
- The response outcome for each female crab is her number of satellites.
- We have several factors (including the female crab's color, spine condition, weight, and carapace width) which may influence the presence/absence of satellite males.
- The data is called hcrabs in the R package rsq.

■ Let's fit the Poisson regression model to the data. In vector form, we have

where is the number of satellites for female crab, and contains the intercept and female crab's

- color;
- spine condition;
- weight; and
- carapace width.
- Suppose we specify a normal prior for
- Can you write down the posterior for ? Can you sample directly from it?

- We can use Metropolis to generate samples from the posterior.
- First, we need a "symmetric" proposal density ; a reasonable choice is usually a multivariate normal centered on
- What about the variance of the proposal density? We can use the variance of the ols estimate, that is,
 , which we can scale using , to tune the acceptance ratio.
- Here, is calculated as the sample variance of , for some small constant , to avoid problems when .
- So we have
- Finally, since we do not have any information apriori about , let's set the prior for it to be
 .

■ The Metropolis algorithm for this mode	el is	S
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1. Given a current , generate a candidate value

.

2. Compute the acceptance ratio

3. Sample

and set



IN-CLASS ANALYSIS: MOVE TO THE R SCRIPT HERE.



- Based on the results from the R script, we have that the expected count of male satellites
 - decreases by a multiplicative factor of for the group with color=4 (medium dark) in comparison to baseline group with color=2 (medium light). That is, a 39% decrease.
 - increases by a multiplicative factor of for the group with spine=3 (both worn or broken) in comparison to baseline group with spine=1 (both good). That is, an 8% increase.
- Both width and weight increases the expected count of male satellites.
- Take a look at the Cls to quantify uncertainty.
- We can actually do a better job with model selection but I leave that to you!!



- Gibbs sampling and the Metropolis algorithm are special cases of the Metropolis-Hastings algorithm.
- The Metropolis-Hastings algorithm is more general in that it allows arbitrary proposal distributions.
- These can be symmetric around the current values, full conditionals, or something else entirely as long as they do not depend on values in our sequence that are previous to the most current values.
- That last point is to ensure the sequence is a Markov chain!
- In terms of how this works, the only real change from before is that now, the acceptance probability should also incorporate the proposal density when it is not symmetric.

Suppose our target distribution is	. The algorithm proceeds as
follows:	

- 1. Given a current draw , propose a new value
- 2. Compute the acceptance ratio

3. Sample and set



and set

for us, then we have	posterior distribution	as is often the case
1. Propose a new value		
2. Compute the accepto	ance ratio	



3. Sample

- Suppose our target distribution is , a bivariate distribution for random variables and .
- For example, could be the joint posterior distribution for and
- Two options:
 - Define one joint proposal density for and if possible; or
 - Define two proposal densities, one for and the other for . That is,
 and .
- First option follows directly from the main algorithm and often works very well when possible. Second option needs a little modification.



1. Update : first, sample . Then,

Compute the acceptance ratio

Sample . Set to if , or set to otherwise.

2. Update : first sample . Then,

Compute the acceptance ratio

Sample . Set to if , or set to otherwise.

- The acceptance ratio looks like what we had before except with an additional factor.
- That factor is the ratio of the probability of generating the current value from the proposed to the probability of generating the proposed value from the current (ratio is equal to one for symmetric proposal see homework!).
- Also, it is often the case that full conditionals are available for some parameters but not all.
- Very useful trick is to combine Gibbs and Metropolis. We may get to that very briefly next time if we have time.