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Final Paper for Course Advanced Quantitative Methods in Political Science

On Using the Metropolis-Hastings Algorithm for Data Imputation

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1 Introduction

dem support + sensitivity analysis

2 Literature Review

In 2020 Christopher Claassen authors two publications, Claassen (2020a) and Claassen (2020b) where he studies the concept of public support for democracy (PSD).

In the earlier article, Claassen (2020a), he investigates the impact of changes in public democratic support on measures of actual democracy. In particular, he has the following three hypothesis: H1: PSD is positively associated with subsequent change in democracy regardless of the initial level of democracy, H1-dem: PSD in already-existing democracies is positively associated with subsequent change in democracy, and H1-aut: PSD in autocracies is positively associated with subsequent change in democracy. Claassen finds evidence for H1-dem, mixed evidence for H1 and no evidence for H1-aut.

In the subsequent article, Claassen (2020b), he analyzes the drivers of PSD. There, he reframes PSD as democratic mood. Claassen has the following hypotheses: H2: Increases in democracy have a negative effect on PSD, H2-elec: increases in electoral democracy have negative effect on PSD, and H2-min: increases in minoritarian democracy have a negative effect on PSD. The evidence support H2 and H2-min but not H2-elec.

Both publications use panel data for PSD in many countries and measured by multiple items. However, the respective datasets display many gaps. To solve this problem, Claassen describes a method to estimate the data in the first publication of the three-part series, Claassen (2019).

The two aims of this paper are (i) to check how stable the estimated data is, and (ii) how sensitive the results are with respect to changes in its hyperparameters. The next section explains the background and the section thereafter describes Claassen's model for PSD and the estimation method.

3 Background

3.1 Introduction

The Metropolis-Hastings (MH) algorithm is a method for sampling data points from a probability distribution from which direct sampling is difficult. It places among the top 10 algorithms with the greatest influence on science and engineering in the 20th century (Beichl and Sullivan 2000). The MH algorithm belongs to the class of Markov chain Monte Carlo (MCMC) methods. In my explanation I assume prior knowledge on Monte Carlo sampling. However, I will describe the basics of Markov Chains. This section is structured as follows. First, I motivate the usage of the MH algorithm. Second, I explain the basics of Markov Chains. Third, I derive the algorithm and make clear why it works.

3.2 Motivation

One main application for the MH algorithm is Bayesian inference. Specifically, we want to estimate parameters θ of some probabilistic model f. We have only limited prior knowledge of the distribution of θ , $p(\theta)$, and we have a likelihood sample of f given the unknown θ , namely $p(X|\theta)$. The goal is to estimate the posterior distribution of θ , $p(\theta|X)$, given all information that we have. In practice, we do not have a formal definition of the likelihood but only observations. Therefore, we can only approximate the posterior by numerical integration, i.e., we need to sample many points from the posterior to describe it. We can then use the posterior sample to estimate θ with the maximum a posteriori probability estimate.

Claassen (2019) uses the MH algorithm to impute gaps in a panel data set. His approach consists of four steps: First, assume a data generating process f parameterized by θ . Second, provide the algorithm with the incomplete data X as likelihood and select priors for θ to obtain the posterior distribution $p(\theta|X) = \frac{p(X|\theta)p(\theta)}{P(X)} \propto p(X|\theta)p(\theta)$. Third, use the values for θ with the highest posterior probability as estimates for θ . Finally, insert these estimates into the assumed probabilistic, data generating model f and use it to sample the missing

data.

In general and abstracting from Bayesian inference, the MH algorithm generates a sample of random states according to the desired probability distribution P(X). For this purpose, the algorithm employs a Markov process that converges to a unique stationary distribution $\pi(x)$ with $\pi(x) = P(X)$. This distribution can then be used for further steps as previously described. The next section explains the conceptual basics.

3.3 Markov Chains

A Markov chain $(X_t)_{t\in\mathbb{N}}$ is a stochastic process (over time) with the property that the probability of the realization in the next period depends solely on the realization in the current state and not the complete history. This is called the Markov property. Because Markov chains with a countable, or discrete, state space are much more accessible than their continuous variant, in this chapter we will look at the discrete case. Formally, the Markov property writes

$$P(X_{t+1}|X_t, X_{t-1}, ..., X_0) = P(X_{t+1}|X_t).$$
(1)

Under some conditions, the stochastic process described by a Markov chain converges to a time-invariant probability distribution, i.e. $P(X_{t+k}|X_{t+k-1}) = P(X_t|X_{t-1}), \forall k > 0$. The crucial step for understanding the MH is to see how it samples a Markov Chain that is certain to converge to a stable posterior distribution. Before exploring how the MH algorithm achieves this result, however, it is necessary to understand its conditions conceptually. To this end, we will use the example depicted by the following graph in Figure 1 that shows the intertemporal transition probabilities between three states representing random events.

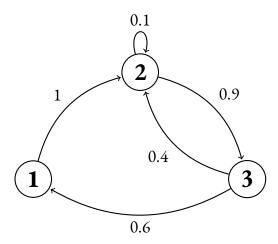


Figure 1: Transition Graph for Markov Chain with 3 states.

This transition graph can be summarized by the $n \times n$ transition matrix T where each element (i, j) represents the probability of moving from state i in period t to state k in period t+1, and where n represents the number of states, i.e $T_{i,j}=P(X_{t+1}=j|X_t=i)$. For our example, we have

$$T = \begin{pmatrix} 0 & 1 & 0 \\ 0 & 0.1 & 0.9 \\ 0.6 & 0.4 & 0 \end{pmatrix}. \tag{2}$$

3.3.1 Limit Distribution

As touched upon in the previous subsection, interesting questions can be what the probabilities of each state $j \in \{1, ..., s\}$ are after a finite number or infinitely many steps. For this purpose let $\pi_t(j) = P(X_t = j)$ denote the probability of being in state j in period t. Of course, the probabilities in t > 0 depend on the probabilities for the the initial state π_0 . We can use the law of total probability to calculate the probability of each state for the next period t = 1 by

$$P(X_1 = j) = \sum_{i=1}^{3} P(X_1 = j | X_0 = i) \pi_0(i).$$
 (3)

I.e., to compute the probability of being in state j in t = 1, for each initial state i, we multiply its probability $\pi_0(i)$ by the probability of moving from i to state j.

This is equivalent to $\pi_1 = \pi_0 T$ in vector notation. Further, we can compute the distributions in an arbitrary future period by repeating the matrix multiplication, e.g, $\pi_2 = \pi_0 T T$, or in general, $\pi_t = \pi_0 T^t$.

Now we are ready to define the limit distribution that describes the probability distribution after infinitely many periods by

$$\pi_{\infty} = \lim_{t \to \infty} \pi_t = \lim_{t \to \infty} \pi_0 T^t. \tag{4}$$

We can further ask two additional important questions. First, does a limit distribution exist? And second, is it unique, or in other word, do we have the same limit distribution independent from the realization of the initial state X_0 ? In our example, there does not only exist a limit distribution with $\pi_{\infty} = (0.2, 0.4, 0.4)$, it is even unique regardless of start distribution π_0 . This means that independent of the start state, the probability of each state converges to the same number. For the context of the MH algorithm, this is an important property because we always want to compute the same estimates for our parameters θ , regardless of the starting values of our simulation. In the next section, we introduce and simplify conditions that guarantee a unique limit distribution.

3.3.2 Irreducibility, Periodicity and Stationarity

Definition 3.1. A Markov chain is called *irreducible* if each state is reachable from any other state in a finite number of steps.

Figure 2 shows a Markov chain represented by a bipartite graph. This graph is composed by two times the graph in Figure 1. Obviously, this chain is not irreducible because the initial state impacts all future distributions. More precisely, starting in one subgraph sets the probability of reaching states in the other subgraph to zero. We see that a Markov Chain is only irreducible if there is at least an indirect link between every pair of states. We also observe that if the Markov Chain is not irreducible there can be no limit distribution.

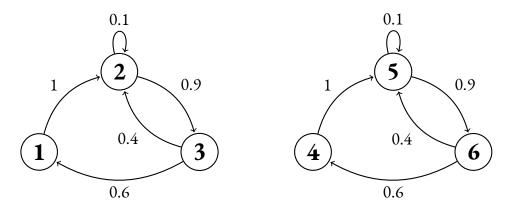


Figure 2: Transition Graph for Irreducible Markov Chain.

Definition 3.2. A state i has a period k if the greatest common denominator of possible revisits is k. A Markov chain is *aperiodic* if the period of all its states is 1.

Consider the five-state Markov chain in Figure 3 as an illustration for the above definition and suppose we start in state 1. Observe that, independent of the random draw for next period, we will arrive again in state 1 after two or four steps. Therefore, state 1 has a period of 2. If a state is revisited in random rather than a fixed time period then the state has period 1. This is automatically the case if a state has a positive edge with itself.

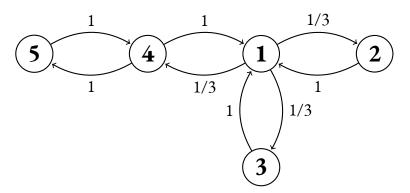


Figure 3: Markov Chain with 2-periodic State 1

Definition 3.3. π^* is the *stationary distribution* of a Markov Chain with Transition matrix T if $\pi^* = \pi^* T$ and π^* is a probability vector.

Verbally, this means that the probability distribution π^* does not change anymore over time. If π^* is also unique, then π^* is our aim, the limit distribution introduces in section 1.3.1, i.e., $\pi^* = \pi_{\infty}$.

These three definitions are enough to understand the next fundamental theorem.

3.3.3 The Fundamental Theorem of Markov Chains

The next theorem defines formally the condition when a Markov Chain converges to a unique distribution, i.e. the limit distribution.

Theorem 3.1. (Fundamental Theorem of Markov Chains) If a Markov chain is irreducible and aperiodic (called ergodic) then it has a stationary distribution π^* that is unique $(\lim_{t\to\infty} P(X_t=i)=\pi_i^*, \forall i)$.

Therefore, if we want to construct a stable distribution P(X) via Markov chains, we need to ensure that it is irreducible and aperiodic with stationary distribution $\pi^* = P(X)$. In the next subsection, we substitute the stationarity condition by a stronger one before we finally derive the MH algorithm.

3.3.4 Reversibility

Definition 3.4. A Markov chain is *reversible* if there is a probability distribution π over its states such that $\pi(i)T_{ij} = \pi(j)T_{j,i}, \forall i, j$ (reversibility condition).

Theorem 3.2. A sufficient condition for distribution π^* to be a stationary distribution of a Markov chain with transition matrix T is that it fullfills the reversibility condition.

Proof.
$$\sum_{i} \pi(i) T_{i,j} = \sum_{i} \pi(j) T_{j,i} = \pi(j) \sum_{i} T_{j,i} = \pi(j) \implies \pi T = \pi$$

Reversibility is a stronger condition than stationarity because it requires that the probability flux from i to j is equal to the one from j to i for each possible pair of states. Recall, that stationarity only requires that the probability flux to one state is equal on aggregate and not that it is symmetric between each pair of states over time. Therefore, if we want to achieve a stationary distribution it is enough to ensure that it is reversible.

3.4 The Algorithm

Recall that we want to generate a sample of a desired distribution P(X). For this purpose, we use a Markov process that is uniquely defined by its transition probabilities $P(X_{t+1}|X)$ with limit distribution π so that $\pi = P(X)$. As explained in the previous section, a Markov process has a limit distribution if each transition $X_t \to X_{t+1}$ is reversible and if the stationary distribution π is ergodic. With the MH algorithm, we construct such a Markov process with stationary distribution $\pi = P(X)$. The derivation starts with another way of writing reversibility¹:

$$P(x'|x)P(x) = P(x|x')P(x') \iff \frac{P(x'|x)}{P(x|x')} = \frac{P(x')}{P(x)}$$
(5)

The main idea is to separate transition P(x'|x) in two steps: the proposal step and the acceptance-or-rejection step. Let g(x') be the proposal distribution, i.e., the conditional probability of proposing state x' given x. And let A(x'|x) be the probability of accepting proposed state X'. Formally, we have P(x'|x) = g(x'|x)A(x'|x'). Inserting this in Equation (5) gives

$$\frac{P(x')}{P(x)} = \frac{g(x'|x)A(x',x)}{g(x|x')A(x',x)} \iff \frac{A(x',x)}{A(x,x')} = \frac{P(x')}{P(x)}\frac{g(x|x')}{g(x'|x)}.$$
 (6)

The following choice, termed the Metropolis choice, is commonly used as an acceptance ratio for sampling x' from P(x') that fulfills the above reversibility condition:

$$A(x',x) = \min\left(1, \frac{P(x')}{P(x)} \frac{g(x|x')}{g(x'|x)}\right) \tag{7}$$

Note that the minimizer in A(x', x) enforces that the probability is below 1. The MH algorithm writes as follows:

¹We simplify our notation by using x' and x instead of X_{t+1} and X_t .

Algorithm 1 Metropolis-Hastings algorithm

Obviously, the construction of the acceptance ratio ensures reversibility. Ergodicity is ensured by the random nature with which we accept proposed states: First, the chain is irreducible because each state is reachable from any other state with positive probability at every single step. Second, for each state x, P(x'=x) is always positive and therefore the Markov chain is aperiodic.

In a general setting, the choice for transition distribution g(x'|x) and the number of iterations until the limit distribution is reached are unclear. These two choices are the hyperparameters of the MH algorithm. In the Bayesian inference application in the article series staring from Claassen (2019), additional choices are the prior distribution $p(\theta)$ and the model choice f.

4 Model and Estimation

Explain Method in 2018b

5 Discussion

6 Conclusion

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