

# MCMC techniques

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- 5.1 Markov chains (R12.1)**
- 5.2 Metropolis-Hastings (R12.2)**
- 5.3 Gibbs sampling (R12.3)**

## 5.1 Markov chains

A **discrete time Markov chain**  $\{X_k\}_k$  is a sequence of rvs such that the distribution of  $X_k$  depends only on  $X_{k-1}$  and not on the previous rvs.

The Markov chain is **homogeneous** if the conditional distribution does not depend on time point. In such a case, the probabilities  $P_{i,j} = P(X_k = j | X_{k-1} = i)$  are called **transition probabilities**.

A pmf  $\pi$  is **stationary distribution** of the Markov chain if for every state  $\pi(j) = \sum P_{i,j}\pi(i)$ .

**Ergodic** Markov chains have a unique stationary distribution and it will be reached as a limit distribution from any initial distribution.

**MCMC** methods consist in generating ergodic Markov chains whose limit distribution is a goal distribution. They do NOT produce sequences of independent observations!

## 5.2 Metropolis-Hastings

Our goal is to simulate from dmf  $f$  and we use an instrumental conditional dmf  $g(\cdot|\cdot)$  from which it is simple to simulate. The support of  $g$  must contain that of  $f$ .

- 1 Set  $X_0$  such that  $f(X_0) > 0$  and  $k = 0$ .
- 2 Set  $k = k + 1$ . Generate  $Y_k \sim g(\cdot|X_{k-1})$  and  $U$  random number in  $(0, 1)$  independent.
- 3 If  $U \leq \alpha(X_{k-1}, Y_k)$ , set  $X_k = Y_k$ , otherwise  $X_k = X_{k-1}$ .
- 4 If  $k < n$  go to Step 2., otherwise return  $X_0, \dots, X_n$ .

$$\alpha(x, y) = \min \left\{ \frac{f(y)g(x|y)}{f(x)g(y|x)}, 1 \right\}$$

# Metropolis-Hastings (Poisson)

In order to simulate a Poisson rv, we take as instrumental probability mass function (given  $x \neq 0$ ), the one that assesses probability  $1/2$  to  $x - 1$  and probability  $1/2$  to  $x + 1$ . If  $x = 0$ , then the probability of 0 is  $1/2$  and the probability of 1 is  $1/2$

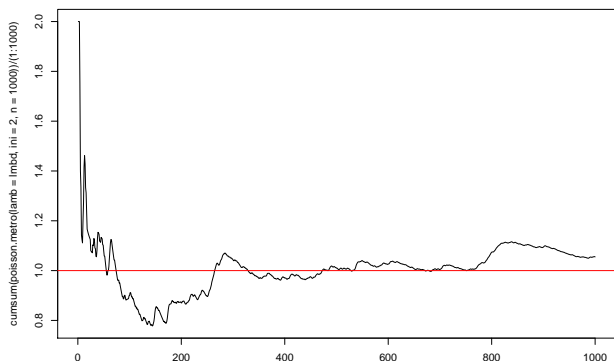
```
rinst <- function(x) {  
  if(x==0) return(sample(1,x=c(0,1)))  
  else return(sample(1,x=c(x-1,x+1)))  
}
```

# Metropolis-Hastings (Poisson)

```
poisson.metro=function(lamb,ini,n){
  x <- c(ini)
  y <- rinst(ini)
  for(k in 2:n) {
    u <- runif(1)
    alpha<-lamb^y/factorial(y)/(lamb^x[k-1]/factorial(x[k-1]))
    if(u<=alpha) x <- c(x,y)
    else x <- c(x,x[k-1])
    y <- rinst(x[k])
  }
  return(x)
}
```

# Metropolis-Hastings (Poisson)

```
lmbd <- 1
plot(cumsum(poisson.metro(lamb=lmbd,ini=2,n=1000))/(1:1000),
     type="l")
abline(h=lmbd,col="red")
```



## 5.3 Gibbs sampling

Consider a random vector  $\mathbf{X} = (X^{(1)}, \dots, X^{(d)})$  with joint dmf  $f$  difficult to simulate from, but whose conditional densities of each component given the remaining components  $f_i(\cdot | X_k^{(1)}, \dots, X_k^{(i-1)}, X_k^{(i+1)}, \dots, X_k^{(d)})$  are easy to simulate from.

The **Gibbs sampler** is Metropolis-Hastings algorithm with  $g(\cdot | \mathbf{X})$  having marginal densities as above. In such a case,  $f(\mathbf{y})g(\mathbf{x} | \mathbf{y}) = f(\mathbf{x})f(\mathbf{y})$ , so all candidates will be accepted.

- 1 Set  $\mathbf{X}_0$  such that  $f(\mathbf{X}_0) > 0$  and  $k = 0$ .
- 2 Set  $k = k + 1$ . For  $i = 1, \dots, d$ , generate  $X_k^{(i)}$  from  $f_i(\cdot | X_k^{(1)}, \dots, X_k^{(i-1)}, X_k^{(i+1)}, \dots, X_k^{(d)})$ .
- 3 If  $k < n$  go to Step 2., otherwise return  $\mathbf{X}_0, \dots, \mathbf{X}_n$ .



## Gibbs sampling (Example)

If the random vector  $(X, Y)^t$  follows a bivariate normal distribution with mean vector  $(0, 0)^t$  and covariance matrix having 1s on the main diagonal and  $\rho$ s elsewhere ( $X$  and  $Y$  are standard normal random variables with correlation  $\rho$ )

$$X \sim N(\rho Y, \sqrt{1 - \rho^2}) \text{ and } Y \sim N(\rho X, \sqrt{1 - \rho^2})$$

## Gibbs sampling (Example)

```
set.seed(1) ; n <- 10^4 ; x <- c(0) ; y <- c(0)
rho <- 0.8 ; sigma <- sqrt(1 - rho^2)
for (i in 2:n) {
  x <- c(x,rnorm(1,mean=rho*y[i-1],sd=sigma))
  y <- c(y,rnorm(1,mean=rho*x[i],sd=sigma))
}
c(mean(x),mean(y))
```

```
## [1] -0.01986475 -0.01208011
```

```
c(sd(x),sd(y))
```

```
## [1] 0.9947670 0.9899634
```

```
cor(x,y)
```

```
## [1] 0.7084101
```

# Gibbs sampling (Example)

```
par(mfrow=c(2,2))  
plot(x[1:100],type="l") ; acf(x)  
s.norm <- rnorm(100) ; plot(s.norm,type="l") ; acf(s.norm)
```

