Advanced Statistical Computing

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Markov Chain Monte Carlo

- The goal is to generates sequence of random samples from an arbitrary probability density function (usually high dimensional),
- The sequence of random samples form a Markov chain,
- The purpose is simulation (Monte Carlo).

What is Monte Carlo



What is Monte Carlo?

- Rely on repeated sampling to study the results of a experiment or study the properties of certain procedure.
 - Often used in complex and uncertain scenarios
 - Difficult to formulate, high correlation.
 - Cheap
 - Take advantage of faster computers
- History
 - John von Neumann, Stanislaw Ulam, Nicholas Metropolis

Applications of Monte Carlo

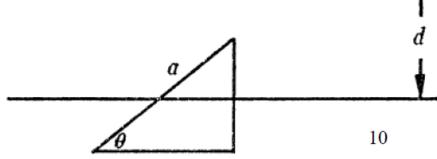
- Optimization
- Numerical integration
- Generate random samples

Buffon's needle

Georges-Louis Leclerc, Comte de Buffon (1707-1788)

Given a needle of length a and an infinite grid of parallel lines with common distance d between them, what is the probability P(E) that a needle, tossed at the grid randomly, will cross one of the parallel lines?





Buffon's needle

• Assume a < d

$$P(E) = \int_0^{\pi} \frac{a \sin \theta d\theta}{\pi d} = (a/\pi d) \int_0^{\pi} \sin \theta d\theta = 2a/\pi d.$$

http://web.student.tuwien.ac.at/~e9527412/buffon.html

Motivation

- Generate *iid* r.v. from high-dimensional arbitrary distributions is extremely difficult.
- Drop the "independent" requirement.
- How about also drop the "identically distributed" requirement?

Markov chain

- Assume a finite state, discrete Markov chain with *N* different states.
- Random process X_n , n = 0,1,2,...

$$x_n \in S = \{1, 2, ..., N\}$$

Markov property,

Markov graph of transiton probabilites between states A, B and C

$$P(X_{n+1} = x \mid X_1 = x_1, X_2 = x_2, ..., X_n = x_n) = P(X_{n+1} = x \mid X_n = x_n)$$

- Time-homogeneous
- Order
 - Future state depends on the past *m* states.

Key parameters

Transition matrix

$$P(X_n = j | X_{n-1} = i) = p(i, j),$$

 $P = \{p(i, j)\}.$

• Initial probability distribution $\pi^{(0)}$

$$\pi^{(n)}(i) = P(x_n = i).$$

• Stationary distribution (invariant/equilibrium)

$$\pi = \pi P$$
.

Reducibility

- A state *j* is accessible from state *i* (written $i \rightarrow j$) if $P(X_n = j \mid X_0 = i) = p_{ij}^{(n)} > 0$.
- A Markov chain is *irreducible* if it is possible to get to any state from any state.

Recurrence

• A state *i* is *transient* if given that we start in state *i*, there is a non-zero probability that we will never return to *i*. State *i* is *recurrent* if it is not transient.

Ergodicity

• A state *i* is *ergodic* if it is aperiodic and positive recurrent. If all states in an irreducible Markov chain are ergodic, the chain is *ergodic*.

Reversible Markov chains

• Consider an ergodic Markov chain that converges to an invariant distribution π . A Markov chain is *reversible* if for all $x, y \in S$,

$$\pi(x)p(x,y) = \pi(y)p(y,x).$$

which is known as the detailed balance equation.

• An ergodic chain in equilibrium and satisfying the detailed balance condition has π as its unique stationary distribution.

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Markov Chain Monte Carlo

- The goal is to generates sequence of random samples from an arbitrary probability density function (usually high dimensional),
- The sequence of random samples form a Markov chain,

in Markov chain, $P \rightarrow \pi$ in MCMC, $\pi \rightarrow P$.

Examples

$$P(X | E, \beta, \sigma^2) \propto \prod_{k=1}^{K} \prod_{E(i)=k} \prod_{j=1}^{M} \left((\sigma_{kj}^2)^{-\frac{1}{2}} e^{-\frac{1}{2\sigma_{kj}^2} (x_{ij} - \beta_{kj})^2} \right)$$

$$P(X \mid E) = \prod_{k=1}^{K} \prod_{i=1}^{M} \iint \prod_{E(i)=k} P(x_{ij} \mid \beta_{kj}, \sigma_{kj}^{2}) p(\beta_{kj} \mid \beta_{0}, \sigma_{kj}^{2}) p(\sigma_{kj}^{2}) d\beta_{kj} d\sigma_{kj}^{2}$$

$$= \prod_{k=1}^{K} \prod_{j=1}^{M} \frac{b^{a}}{\Gamma(a)} \frac{(2\pi)^{\frac{-n_{k}}{2}}}{\sqrt{n_{k}+1}} \frac{\Gamma\left(\frac{n_{k}}{2}+a\right)}{\left(b+\frac{1}{2}\left(\sum_{E(i)=k} x_{ij}^{2}+\beta_{0}^{2}-\frac{\left(\sum_{E(i)=k} x_{ij}+\beta_{0}\right)^{2}}{n_{k}+1}\right)\right)^{\frac{n_{k}}{2}+a}}$$

Bayesian Inference

$$Y = (y_1, \dots, y_n)$$

$$Z = (z_1, \ldots, z_n)$$

Genotype
$$Y = (y_1, ..., y_n)$$

Haplotype $Z = (z_1, ..., z_n)$
Frequency $\Theta = (\theta_1, ..., \theta_m)$

Prior

$$\theta \sim Dirichlet(\beta)$$

$$P(Y, Z, \Theta) = \prod_{i=1}^{n} \theta_{z_{i1}} \theta_{z_{i2}} \prod_{g=1}^{m} \theta_{g}^{\beta_{g}-1}$$

History

• Metropolis, N., Rosenbluth, A. W., Rosenbluth, M. N., Teller, A. H. and Teller, E. (1953).

Equation of state calculations by fast computing machines. *Journal of Chemical Physics*, **21**, 1087–1092.

Metropolis algorithm

- Direct sampling from the target distribution is difficult,
- Generating candidate draws from a proposal distribution,
- These draws then "corrected" so that asymptotically, they can be viewed as random samples from the desired target distribution.

Pseudo code

- Initialize X_0 ,
- Repeat
 - Sample $Y \sim q(x,.)$,
 - Sample $U \sim Uniform (0,1)$,
 - If $U \le \alpha(X,Y)$, set $X_i = y$,
 - Otherwise $X_i = x$.

An informal derivation

- Find $\alpha(X,Y)$:
- Joint density of current Markov chain state and the proposal is $g(x,y) = q(x,y)\pi(x)$
- Suppose q satisfies detail balance $q(x,y) \pi(x) = q(y,x)\pi(y)$
- If q(x,y) $\pi(x) > q(y,x)\pi(y)$, introduce $\alpha(x,y) < 1$ and $\alpha(y,x) = 1$ hence $\alpha(x, y) = \frac{\pi(y)q(y, x)}{\pi(x)q(x, y)}$. • If $q(x,y) \pi(x) > q(y,x)\pi(y)$, ...
- The probability of acceptance is $\alpha(x, y) = \min \left(1, \frac{\pi(y)q(y, x)}{\pi(x)q(x, y)} \right)$.

Metropolis-Hastings Algorithm

- Start with any $X^{(0)} = x_0$, and a "proposal chain" T(x,y)
- Suppose $X^{(t)} = x_t$. At time t+1,
 - Draw $y \sim T(x_t, y)$ (i.e., propose a move for the next step)
 - Compute "goodness ratio"

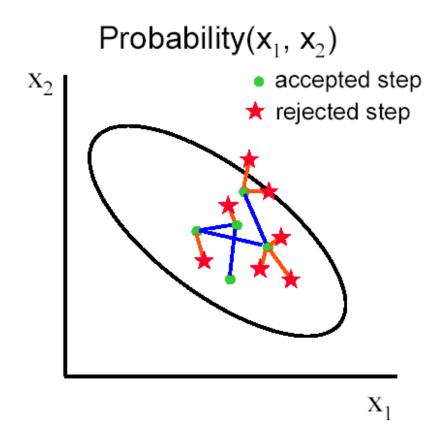
$$r = \frac{\pi(y)T(y, x_t)}{\pi(x_t)T(x_t, y)}$$

Acceptance/Rejection decision: Let

$$X^{(t+1)} = \begin{cases} y, & \text{with } p = \min\{1, r\} \\ x_t, & \text{with } 1 - p \end{cases}$$

Remarks

 Relies only on calculation of target pdf up to a normalizing constant.



Remarks

- How to choose a good proposal function is crucial.
- Sometimes tuning is needed.
 - Rule of thumb: 30% acceptance rate
- Convergence is slow for high dimensional cases.

Illustration of Metropolis-Hastings

- Suppose we try to sample from a bi-variate normal distributions. $N\left(\begin{pmatrix}0\\0\end{pmatrix},\begin{pmatrix}1&\rho\\\rho&1\end{pmatrix}\right)$
- Start from (0, 0)
- Proposed move at each step is a two dimensional random walk $x_{t+1} = x_t + s \cos \theta$

$$x_{t+1} = x_t + s \cos \theta$$

$$y_{t+1} = y_t + s \sin \theta$$

$$with$$

$$s \sim U(0,1)$$

$$\theta \sim U(0,2\pi)$$

Illustration of Metropolis-Hastings

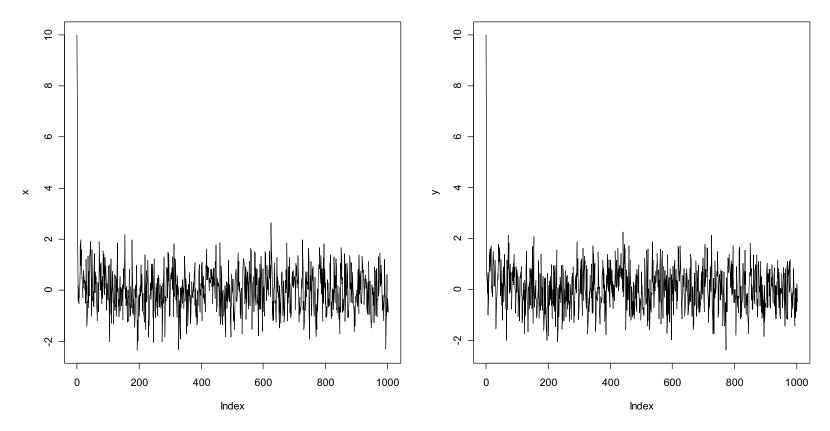
• At each step, calculate $r = \frac{\pi(x_{t+1}, y_{t+1})}{\pi(x_t, y_t)}$

$$T((x_t, y_t), (x_{t+1}, y_{t+1})) = T((x_{t+1}, y_{t+1}), (x_t, y_t)) = 1/\pi^2$$
 since

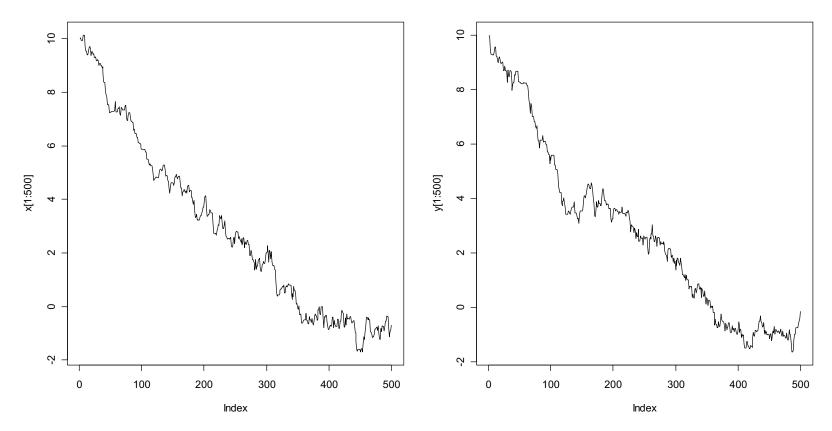
$$r = \frac{\frac{1}{2\pi\sqrt{1-\rho^2}} \exp\left(-\frac{1}{2(1-\rho^2)} \left(x_{t+1}^2 - 2\rho x_{t+1} y_{t+1} + y_{t+1}^2\right)\right)}{\frac{1}{2\pi\sqrt{1-\rho^2}} \exp\left(-\frac{1}{2(1-\rho^2)} \left(x_t^2 - 2\rho x_t y_t + y_t^2\right)\right)}$$

$$= \exp\left(-\frac{1}{2(1-\rho^2)} \left(\left(x_{t+1}^2 - 2\rho x_{t+1} y_{t+1} + y_{t+1}^2\right) - \left(x_t^2 - 2\rho x_t y_t + y_t^2\right)\right)\right)$$

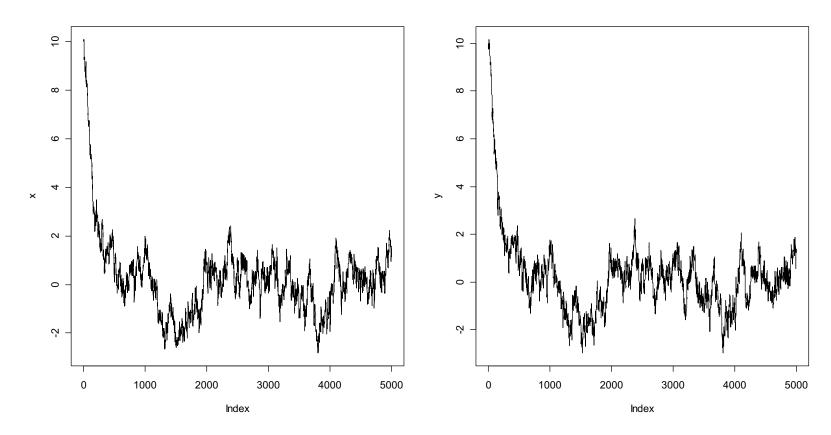
• Trace plot Good



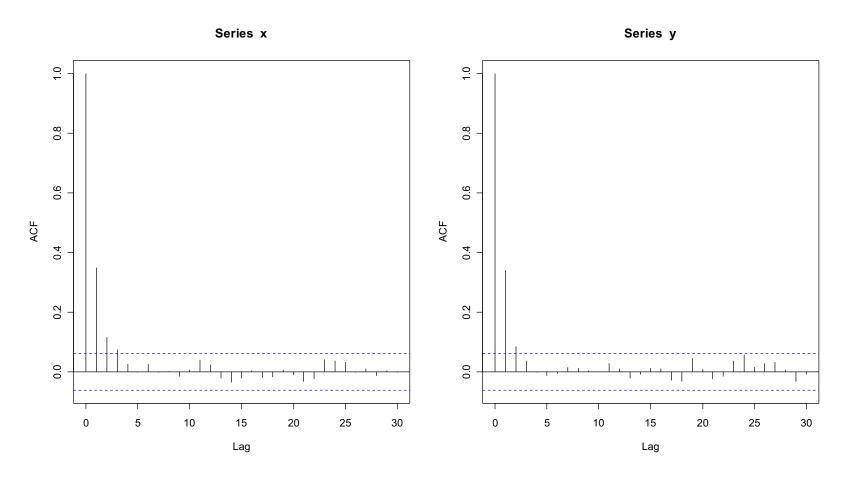
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• Trace plot

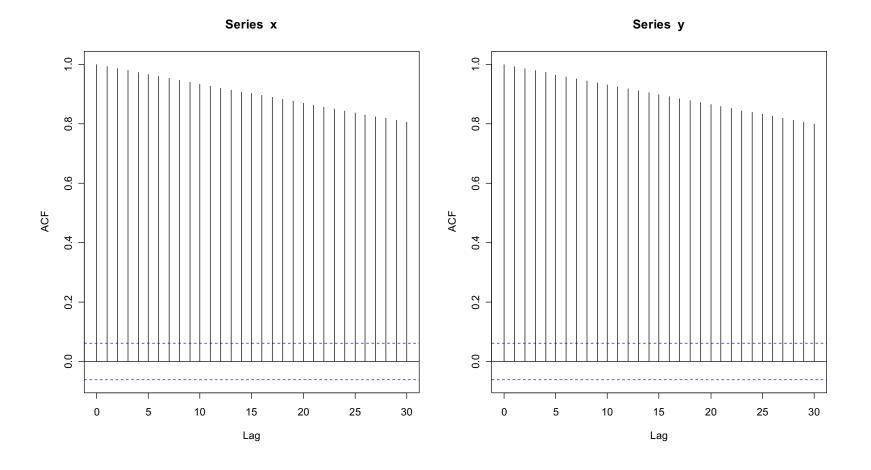


Autocorrelation plot Good



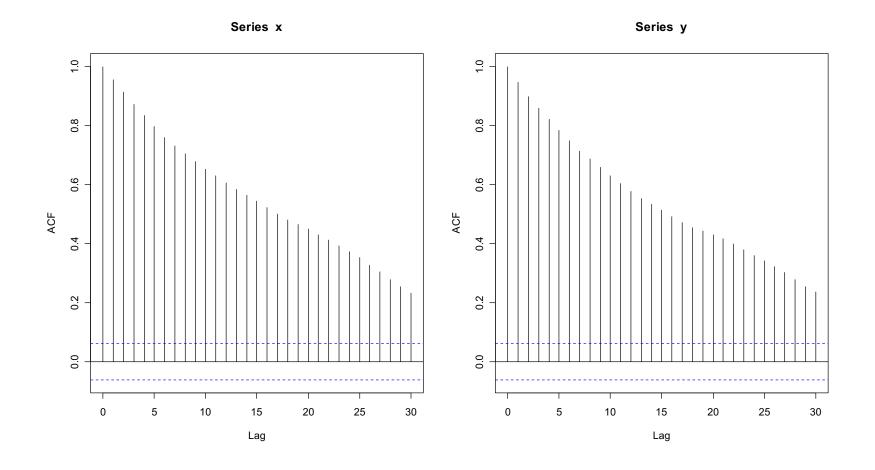
Autocorrelation plot Bad

$$s = 0.5$$



• Autocorrelation plot Okay

$$s = 3.0$$



References

- Metropolis et al. 1953,
- Hastings 1973,
- Tutorial paper:

Chib and Greenberg (1995). Understanding the Metropolis--Hastings Algorithm. *The American Statistician* **49**, 327-335.

Gibbs Sampler

• Purpose: Draw random samples form a joint distribution (high dimensional)

$$x = (x_1, x_2, ..., x_n)$$
 Target $\pi(x)$

Method: Iterative conditional sampling

$$\forall i$$
, draw $\mathbf{x}_i \sim \pi(\mathbf{x}_i \mid \mathbf{x}_{[-i]})$

Illustration of Gibbs Sampler

Suppose the target distribution is:

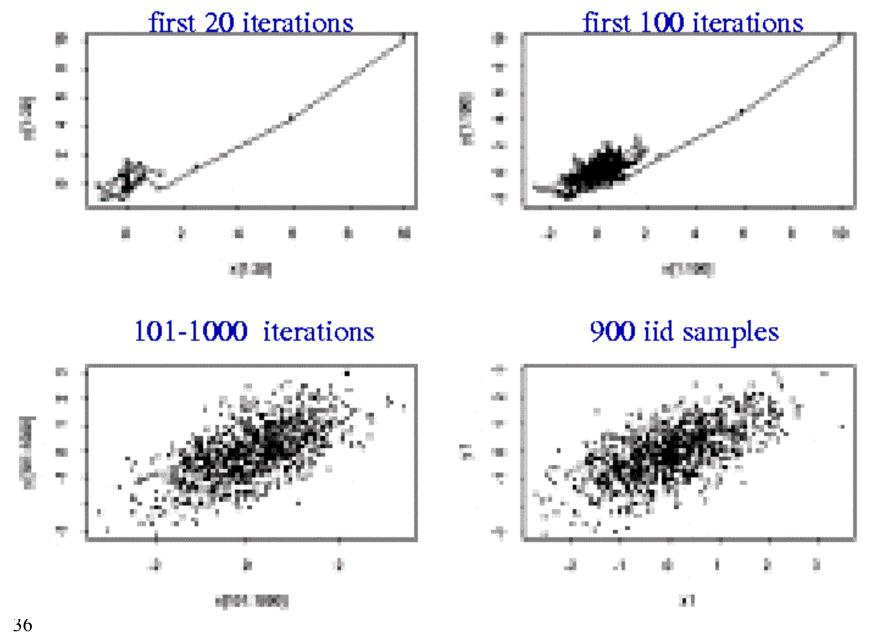
$$(X,Y) \sim N\left(\begin{pmatrix} 0\\ 0 \end{pmatrix}, \begin{pmatrix} 1 & \rho\\ \rho & 1 \end{pmatrix}\right)$$

Gibbs sampler:

$$[X|Y = y] \sim N(\rho y, 1 - \rho^2)$$

$$[Y|X = x] \sim N(\rho x, 1 - \rho^2)$$

Start from, say, (X,Y)=(10,10), we can take a look at the trajectories. We took $\rho=0.6$.



References

- Geman and Geman 1984,
- Gelfand and Smith 1990,
- Tutorial paper:

Casella and George (1992). Explaining the Gibbs sampler. *The American Statistician*, **46**, 167-174.

Remarks

- Gibbs Sampler is a special case of Metropolis-Hastings
- Compare to EM algorithm, Gibbs sampler and Metropolis-Hastings are stochastic procedures
- Verify convergence of the sequence
- Require Burn in
- Use multiple chains

References

- Geman and Geman 1984,
- Gelfand and Smith 1990,
- Tutorial paper:

Casella and George (1992). Explaining the Gibbs sampler. *The American Statistician*, **46**, 167-174.