

- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Bayesian Statistics and Data Analysis Lecture 5

Måns Magnusson Department of Statistics, Uppsala University Thanks to Aki Vehtari, Aalto University



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It's all about expectations

$$E_{p(\theta|y)}[f(\theta)] = \int f(\theta)p(\theta|y)d\theta,$$
 where
$$p(\theta|y) = \frac{p(y|\theta)p(\theta)}{\int p(y|\theta)p(\theta)d\theta}$$



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We can easily evaluate $p(y|\theta)p(\theta)$ for any θ , but the integral $\int p(y|\theta)p(\theta)d\theta$ is usually difficult.



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- Diagnostics
 - vvarm-upConvegernce
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We can use the unnormalized posterior $q(\theta|y) = p(y|\theta)p(\theta)$, for example, in

• Monte Carlo methods which can sample from $p(\theta^{(s)}|y)$ using only $q(\theta^{(s)}|y)$

$$E_{p(\theta|y)}[f(\theta)] \approx \frac{1}{S} \sum_{s=1}^{S} f(\theta^{(s)})$$



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- Diagnostics
 - Warm-up
 - Convegernce
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- Monte Carlo methods we have discussed so far
 - Inverse CDF works for 1D



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 - Convegernce
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 - Convegernce
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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
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 - Convegernce
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- What to do in high dimensions?
 - Markov chain Monte Carlo (Ch 11-12)
 - Laplace, Variational*, EP* (Ch 4, 13*, next course)



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- Diagnostics
 - Warm-up
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 - S_{eff}, MCSE, and autocorrelation

 Andrey Markov proved weak law of large numbers and central limit theorem for certain dependent-random sequences, which were later named Markov chains



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- Andrey Markov proved weak law of large numbers and central limit theorem for certain dependent-random sequences, which were later named Markov chains
- The probability of each event depends only on the state attained in the previous event (or finite number of previous events)

$$p(\theta_t|\theta_{t-1},\theta_{t-2},...)=p(\theta_t|\theta_{t-1})$$



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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 - S_{eff}, MCSE, and autocorrelation

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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
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 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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- Goal in MCMC: Construct a transition distribution with $p(\theta|y)$ as the stationary distribution



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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

• Produce draws $\theta^{(t)}$ given $\theta^{(t-1)}$ from a Markov chain, with stationary distribution $p(\theta|y)$



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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 - + combine sequence of easier Monte Carlo draws to form a Markov chain



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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- Diagnostics
 - Warm-up
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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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 - + asymptotically chain spends the $\alpha\%$ of time where $\alpha\%$ posterior mass is
 - + central limit theorem holds for expectations
 - draws are dependent
 - construction of efficient Markov chains is not always easy



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

• Set of random variables $\theta_1, \theta_2, \ldots$, so that with all values of t, θ_t depends only on the previous $\theta_{(t-1)}$

$$p(\theta_t|\theta_1,\ldots,\theta_{(t-1)}) = p(\theta_t|\theta_{(t-1)})$$



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 - Gibbs sampling
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- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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 - Gibbs sampling
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- Diagnostics
 - Warm up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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- Chain has to be initialized with some starting point θ_0
- Transition distribution $T_t(\theta_t|\theta_{t-1})$ (may depend on t)
- Choose a transition distribution so the stationary distribution of the Markov chain is $p(\theta|y)$



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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- Alternate sampling from conditional distributions
- Basic algorithm, for $j \in \{1,...,J\}$

sample
$$\theta_{j,t}$$
 from $p(\theta_j|\theta_{-j,t-1},y),$ where $\theta_{j,t-1} = (\theta_{1,J},\ldots,\theta_{j-1,t},\theta_{j+1,t-1},\ldots,\theta_{t-1,J})$

• Will converge (in total variation) to $p(\theta|y)$ as $T \to \infty$



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 - Gibbs sampling
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- Diagnostics
 - Warm-up
 - Convegernce
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- 1D sampling (|j| = 1) is generally easy



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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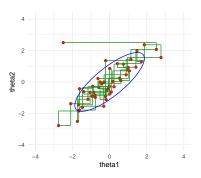
- Will converge (in total variation) to $p(\theta|y)$ as $T \to \infty$
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- Related to the (stochastic) EM algorithm



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- Monte Carlo recap
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 - Gibbs sampling
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- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Gibbs sampling



Draws — Steps of the sampler — 90% HPD

demo



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

 With conditionally conjugate priors, the sampling from the conditional distributions is easy for wide range of models



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- With conditionally conjugate priors, the sampling from the conditional distributions is easy for wide range of models
- BUGS / WinBUGS / OpenBUGS / JAGS



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 Metropolis-Hastings
 - wietropolis-riasting
- Diagnostics
 Warm-up
 - Convegernce
 -
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 Metropolis-Hastings
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- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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- Markov Chain Monte Carlo (MCMC)
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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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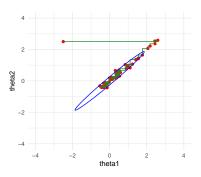
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- BUGS / WinBUGS / OpenBUGS / JAGS
- No algorithm parameters to tune
- For not so easy conditionals, use e.g. inverse-CDF
- Several parameters can be updated in blocks (blocking)
- Slow if parameters are highly dependent in the posterior...



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - ConvegernceConvegernce
 - Seff, MCSE, and autocorrelation

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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Sampling conditional vs joint

- How about sampling θ jointly?
 - e.g. it is easy to sample from multivariate normal



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- Markov Chain Monte Carlo (MCMC)
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- Diagnostics
 - Warm-up
 - Convegernce
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 - S_{eff}, MCSE, and autocorrelation

Sampling conditional vs joint

- How about sampling θ jointly?
 - e.g. it is easy to sample from multivariate normal
- Can we use that to form a Markov chain?



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 - Warm-up
 - Convegernce
 - Convegernce
 Seff, MCSE, and autocorrelation

The Metropolis algorithm

- Algorithm
 - 1. starting point θ^0
 - 2. t = 1, 2, ...
 - (a) pick a proposal θ^* from a proposal distribution $J_t(\theta^*|\theta_{t-1})$.

Proposal distribution has to be symmetric, i.e. $J_t(\theta_a|\theta_b) = J_t(\theta_b|\theta_a)$, for all θ_a, θ_b



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- Monte Carlo recap
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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-upConvegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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 - (b) calculate acceptance ratio

$$r = \frac{p(\theta^*|y)}{p(\theta_{t-1}|y)}$$



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$$\begin{aligned} r &= \frac{p(\theta^*|y)}{p(\theta_{t-1}|y)} \\ \theta_t &= \begin{cases} \theta^* & \text{with probability min}(r,1) \\ \theta_{t-1} & \text{otherwise} \end{cases}$$



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 - Metropolis-Hastings
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 - Warm-up
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 - Convegernce
 S_{eff}, MCSE, and autocorrelation

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ie, if $p(\theta^*|y) > p(\theta_{t-1}|y)$ accept the proposal always and otherwise accept the proposal with probability r



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- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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$$\begin{aligned} r &= \frac{\rho(\theta^*|y)}{\rho(\theta_{t-1}|y)} \\ \theta_t &= \begin{cases} \theta^* & \text{with probability min}(r,1) \\ \theta_{t-1} & \text{otherwise} \end{cases}$$

• rejection of a proposal increments the time *t* also by one ie, the new state is the same as previous



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 - Warm-up
 - Convegernce
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 - S_{eff}, MCSE, and autocorrelation

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- step c is executed by generating a random number from $\mathcal{U}(0,1)$



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- Markov Chain Monte Carlo (MCMC)
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- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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The Metropolis algorithm

- Algorithm
 - 1. starting point θ^0
 - 2. t = 1, 2, ...
 - (a) pick a proposal θ^* from a proposal distribution $J_t(\theta^*|\theta_{t-1})$. Proposal distribution has to be symmetric, i.e. $J_t(\theta_a|\theta_b) = J_t(\theta_b|\theta_a)$, for all θ_a, θ_b
 - (b) calculate acceptance ratio

$$\begin{aligned} r &= \frac{p(\theta^*|y)}{p(\theta_{t-1}|y)} \\ \theta_t &= \begin{cases} \theta^* & \text{with probability min}(r,1) \\ \theta_{t-1} & \text{otherwise} \end{cases}$$

- rejection of a proposal increments the time t also by one ie, the new state is the same as previous
- ullet step c is executed by generating a random number from $\mathcal{U}(0,1)$
- $p(\theta^*|y)$ and $p(\theta_{t-1}|y)$ have the same normalization terms, and thus instead of $p(\cdot|y)$, unnormalized $q(\cdot|y)$ can be used, as the normalization terms cancel out!



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 - Gibbs sampling
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- Diagnostics
 - Warm-up
 - Convegernce
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Metropolis algorithm

- Example: one bivariate observation (y_1, y_2)
 - bivariate normal distribution with unknown mean and known covariance

$$\begin{pmatrix} \theta_1 \\ \theta_2 \end{pmatrix} \middle| \ y \sim \mathcal{N} \left(\begin{pmatrix} y_1 \\ y_2 \end{pmatrix}, \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix} \right)$$

• proposal distribution $J_t(\theta^*|\theta_{t-1}) = \mathcal{N}(\theta^*|\theta_{t-1}, \sigma_p^2)$

demo



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

 Intuitively more draws from the higher density areas as jumps to higher density are always accepted and only some of the jumps to the lower density are accepted



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 Warm-up
 - Convegernce
 -
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- Intuitively more draws from the higher density areas as jumps to higher density are always accepted and only some of the jumps to the lower density are accepted
- Theoretically
 - Prove that simulated series is a Markov chain which has unique stationary distribution
 - 2. Prove that this stationary distribution is the desired target distribution



- Monte Carlo recap
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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- Prove that simulated series is a Markov chain which has unique stationary distribution
 - a) irreducible
 - b) aperiodic

c) recurrent / not transient



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- Prove that simulated series is a Markov chain which has unique stationary distribution
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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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 - holds for a random walk on any proper distribution (except for trivial exceptions)
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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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 - = aperiodic (return times are not periodic)
 - holds for a random walk on any proper distribution (except for trivial exceptions)
 - c) recurrent / not transient
 - = probability to return to a state i is 1 as $T \to \infty$
 - holds for a random walk on any proper distribution (except for trivial exceptions)



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- 2. Prove that this stationary distribution is the desired target distribution $p(\theta|y)$
 - consider starting algorithm at time t-1 with a draw $heta_{t-1} \sim p(heta|y)$



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Convegernce
 - Convegernce
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 - the unconditional probability density of a transition from θ_a to θ_b is

$$p(\theta_{t-1} = \theta_a, \theta_t = \theta_b) = p(\theta_a|y)J_t(\theta_b|\theta_a),$$



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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 - S_{eff}, MCSE, and autocorrelation

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$$p(\theta_{t-1} = \theta_a, \theta_t = \theta_b) = p(\theta_a|y)J_t(\theta_b|\theta_a),$$

- the unconditional probability density of a transition from θ_b to θ_a is

$$p(\theta_t = \theta_a, \theta_{t-1} = \theta_b) = p(\theta_b|y)J_t(\theta_a|\theta_b)\left(\frac{p(\theta_a|y)}{p(\theta_b|y)}\right)$$
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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-un
 - Convegernce
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 - S_{eff}, MCSE, and autocorrelation

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which is the same as the probability of transition from θ_a to θ_b , since we have required that $J_t(\cdot|\cdot)$ is symmetric



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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$$= p(\theta_a|y)J_t(\theta_a|\theta_b),$$

which is the same as the probability of transition from θ_a to θ_b , since we have required that $J_t(\cdot|\cdot)$ is symmetric

- since their joint distribution is symmetric, θ_t and θ_{t-1} have the same marginal distributions, and so $p(\theta|y)$ is the stationary distribution of the Markov chain of θ



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 Convegernce
 - -----g-----
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- Generalization of Metropolis algorithm for non-symmetric proposal distributions
 - acceptance ratio includes ratio of proposal distributions

$$r = \frac{p(\theta^*|y)/J_t(\theta^*|\theta_{t-1})}{p(\theta_{t-1}|y)/J_t(\theta_{t-1}|\theta^*)}$$



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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$$r = \frac{p(\theta^*|y)/J_t(\theta^*|\theta_{t-1})}{p(\theta_{t-1}|y)/J_t(\theta_{t-1}|\theta^*)} = \frac{p(\theta^*|y)J_t(\theta_{t-1}|\theta^*)}{p(\theta_{t-1}|y)J_t(\theta^*|\theta_{t-1})}$$



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 Seff, MCSE, and autocorrelation

- Ideal proposal distribution is the distribution itself
 - $J(\theta^*|\theta) \equiv p(\theta^*|y)$ for all θ
 - acceptance probability is 1
 - independent draws
 - not usually feasible



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 Metropolis-Hastings
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- Diagnostics
 - Warm-upConvegernce
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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 Metropolis-Hastings
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- Diagnostics
 - Warm-up
 - Convegernce
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- After the proposal distribution shape has been selected, it is important to select the scale
 - small scale
 - ightarrow many steps accepted, but the chain moves slowly due to small steps
 - big scale
 - ightarrow long steps proposed, but many of those rejected and again chain moves slowly

demo



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 - Gibbs sampling
 Metropolis-Hastings
 -
- Diagnostics
 - Warm-up
 - Convegernce
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Generic rule for rejection rate is 60-90%



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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Gibbs sampling as a special case

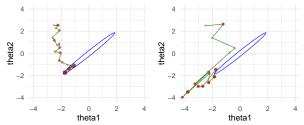
- Specific case of Metropolis-Hastings algorithm
 - single updated (or blocked)
 - proposal distribution is the conditional distribution
 - \rightarrow proposal and target distributions are same
 - ightarrow acceptance probability is 1



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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
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Metropolis

- Usually doesn't scale well to high dimensions
 - if the shape doesn't match the whole distribution, the efficiency drops



- Draws—Steps of the sampler—90% HPI
- Draws-Steps of the sampler-90% HPI



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- Diagnostics
 - Warm-up
 - Convegernce
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Warm-up and convergence diagnostics

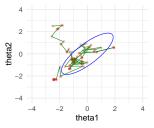
• Asymptotically chain spends the $\alpha\%$ of time where $\alpha\%$ posterior mass is



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
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- Diagnostics
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Warm-up and convergence diagnostics

- Asymptotically chain spends the $\alpha\%$ of time where $\alpha\%$ posterior mass is
 - but in finite time the initial part of the chain may be non-representative and lower error of the estimate can be obtained by throwing it away



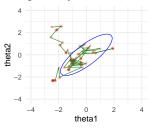
Draws—Steps of the sampler—90% HP



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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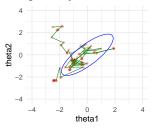
- Draws—Steps of the sampler—90% HPI
- Warm-up = remove draws from the beginning of the chain
 - warm-up may include also phase for adapting algorithm parameters



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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
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 - Convegernce
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Warm-up and convergence diagnostics

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- Draws—Steps of the sampler—90% HPI
- Warm-up = remove draws from the beginning of the chain
 - warm-up may include also phase for adapting algorithm parameters
- Convergence diagnostics
 - Do we get samples from the target distribution?



- Monte Carlo recap
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- Diagnostics
 - Warm-up
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 - Convegernce
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MCMC draws are dependent

Monte Carlo estimates still valid (central limit theorem holds)

$$E_{p(\theta|y)}[f(\theta)] \approx \frac{1}{S} \sum_{s=1}^{S} f(\theta^{(s)})$$

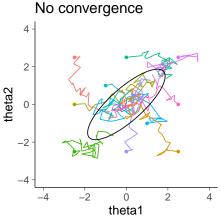
- Estimation of Monte Carlo error is more difficult
 - evaluation of effective sample size



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Assesing convergence: Several chains

- Use of several chains make convergence diagnostics easier
- Start chains from different starting points preferably overdispersed

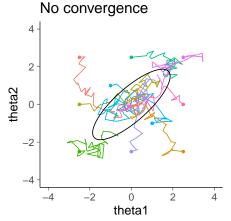




- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 S_{eff}, MCSE, and autocorrelation

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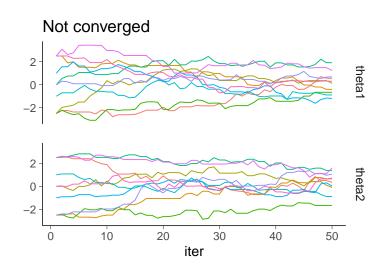


 Remove draws from the beginning of the chains and run chains long enough so that it is not possible to distinguish where each chain started and the chains are well mixed



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - vvarm-u
 - Convegernce
 - Convegernce
 Seff, MCSE, and autocorrelation

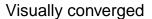
Several chains

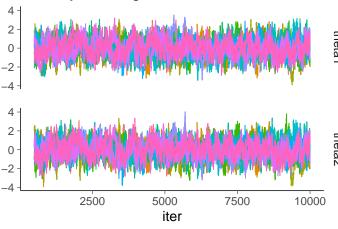




- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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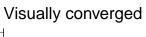


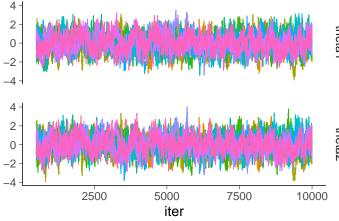




- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Several chains





Visual convergence check is not sufficient



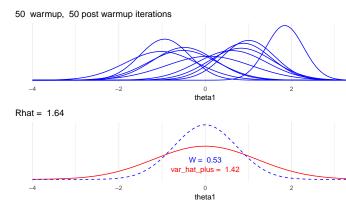
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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

- BDA3: \hat{R} aka potential scale reduction factor (PSRF)
- Compare means and variances of the chains



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

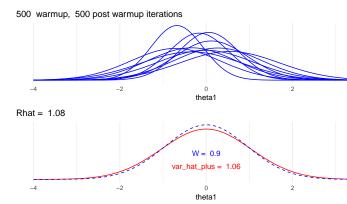
- BDA3: \widehat{R} aka potential scale reduction factor (PSRF)
- Compare means and variances of the chains
 W = within chain variance estimate
 var_hat_plus = total variance estimate





- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
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 - Warm-upConvegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

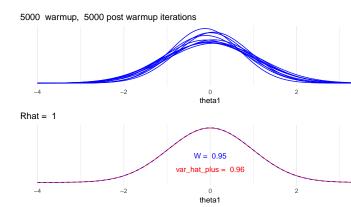
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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



ullet M chains, each having N draws (with new R-hat notation)



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 Seff, MCSE, and autocorrelation



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- M chains, each having N draws (with new R-hat notation)
- Within chains variance W

$$W = rac{1}{M} \sum_{m=1}^{M} s_m^2$$
, where $s_m^2 = rac{1}{N-1} \sum_{n=1}^{N} (heta_{nm} - ar{ heta}_{.m})^2$



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



- *M* chains, each having *N* draws (with new R-hat notation)
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Between chains variance B

$$B = \frac{N}{M-1} \sum_{m=1}^{M} (\bar{\theta}_{.m} - \bar{\theta}_{..})^2,$$

where
$$\bar{\theta}_{.m} = \frac{1}{N} \sum_{n=1}^{N} \theta_{nm}$$
, $\bar{\theta}_{..} = \frac{1}{M} \sum_{m=1}^{M} \bar{\theta}_{.m}$



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



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$$W = \frac{1}{M} \sum_{m=1}^{M} s_m^2$$
, where $s_m^2 = \frac{1}{N-1} \sum_{n=1}^{N} (\theta_{nm} - \bar{\theta}_{.m})^2$

Between chains variance B

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 where $\bar{\theta}_{.m} = \frac{1}{N} \sum_{n=1}^{N} \theta_{nm}, \ \bar{\theta}_{..} = \frac{1}{M} \sum_{n=1}^{M} \bar{\theta}_{.m}$

B/N is variance of the means of the chains



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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- Within chains variance W

$$W = \frac{1}{M} \sum_{m=1}^{M} s_m^2$$
, where $s_m^2 = \frac{1}{N-1} \sum_{n=1}^{N} (\theta_{nm} - \bar{\theta}_{.m})^2$

Between chains variance B

$$B = \frac{N}{M-1} \sum_{m=1}^{M} (\bar{\theta}_{.m} - \bar{\theta}_{..})^{2},$$

where
$$\bar{\theta}_{.m}=rac{1}{N}\sum_{n=1}^{N} heta_{nm},\ \bar{\theta}_{..}=rac{1}{M}\sum_{m=1}^{M}ar{\theta}_{.m}$$

- B/N is variance of the means of the chains
- Estimate total variance $var(\theta|y)$ as a weighted mean of W and B

$$\widehat{\operatorname{var}}^+(\theta|y) = \frac{N-1}{N}W + \frac{1}{N}B$$



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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 - S_{eff}, MCSE, and autocorrelation



• Estimate total variance $\mathrm{var}(\theta|y)$ as a weighted mean of W and B

$$\widehat{\operatorname{var}}^+(\theta|y) = \frac{N-1}{N}W + \frac{1}{N}B$$

• this *overestimates* marginal posterior variance if the starting points are overdispersed



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



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$$\widehat{\operatorname{var}}^+(\theta|y) = \frac{N-1}{N}W + \frac{1}{N}B$$

- this overestimates marginal posterior variance if the starting points are overdispersed
- Given finite N, W underestimates marginal posterior variance
 - single chains have not yet visited all points in the distribution
 - when $N \to \infty$, $E(W) \to var(\theta|y)$



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-upConvegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



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- Given finite N, W underestimates marginal posterior variance
 - single chains have not yet visited all points in the distribution
 - when $N \to \infty$, $E(W) \to var(\theta|y)$
- As $\widehat{\text{var}}^+(\theta|y)$ overestimates and W underestimates, compute

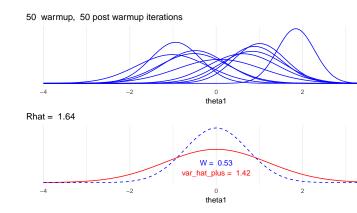
$$\widehat{R} = \sqrt{\frac{\widehat{\text{var}}^+}{W}}$$



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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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- BDA3: \hat{R} aka potential scale reduction factor (PSRF)
- Compare means and variances of the chains
 W = within chain variance estimate
 var_hat_plus = total variance estimate

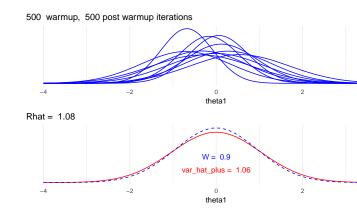




- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



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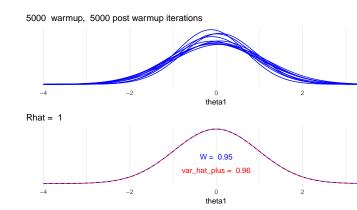




- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



$$\widehat{R} = \sqrt{\frac{\widehat{\text{var}}^+}{W}}$$

- Estimates how much the scale of ψ could reduce if ${\it N} \rightarrow \infty$
- $\widehat{R} \to 1$, when $N \to \infty$
- if \widehat{R} is big (e.g., R > 1.01), keep sampling



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 Metropolis-Hastings
 -
- Diagnostics
 - Warm-upConvegernce
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation



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- $\widehat{R} \to 1$, when $N \to \infty$
- if \widehat{R} is big (e.g., R > 1.01), keep sampling
- If \widehat{R} close to 1, it is still possible that chains have not converged
 - if starting points were not overdispersed
 - distribution far from normal (especially if infinite variance)
 - just by chance when N is finite



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Split- \widehat{R}

- BDA3: split- \widehat{R}
- Examines mixing and stationarity of chains
- To examine stationarity chains are split to two parts
 - after splitting, we have M chains, each having N draws
 - scalar draws θ_{nm} (n = 1, ..., N; m = 1, ..., M)
 - compare means and variances of the split chains



- Monte Carlo recap
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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

• Original \widehat{R} requires that the target distribution has finite mean and variance

Vehtari, Gelman, Simpson, Carpenter, Bürkner (2020). Rank-normalization, folding, and localization: An improved R-hat for assessing convergence of MCMC. Bayesian Analysis, doi:10.1214/20-BA1221.



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- Diagnostics
 - Warm-up
 - Convegernce
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- Original R requires that the target distribution has finite mean and variance
- Rank normalization fixes this and is also more robust given finite but high variance

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 - Gibbs sampling
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- Diagnostics
 - Warm-upConvegernce
 - Convegernce
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- Diagnostics
 Warm-up
 - Convegernce
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 - Warm-up
 - Convegernce
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- Notation updated compared to BDA3

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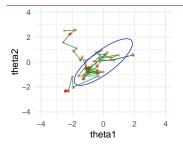
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- Diagnostics
 - Warm-up
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Time series analysis

- Auto correlation function
 - describes the correlation given a certain lag
 - can be used to compare efficiency of MCMC algorithms and parameterizations



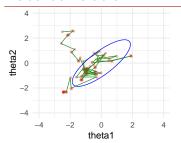
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Draws—Steps of the sampler—90% HPI

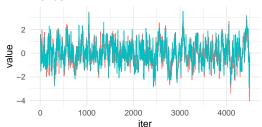


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- Diagnostics
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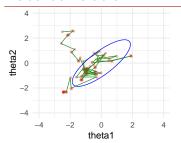
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Trends



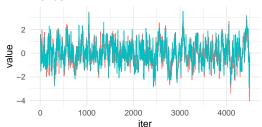


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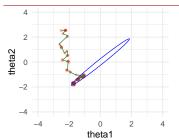
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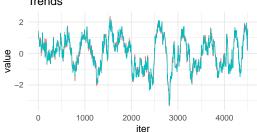


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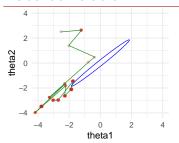
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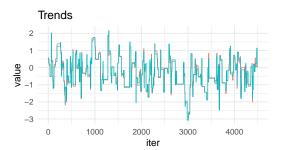




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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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Time series analysis

- Time series analysis can be used to estimate Monte Carlo error in case of MCMC
- For expectation $\bar{\theta}$

$$\operatorname{Var}[\bar{\theta}] = \frac{\sigma_{\theta}^2}{S_{E_{\max}}}$$

where $S_{E_{\max}} = S/\tau$, and τ is sum of autocorrelations



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 $m{ ilde{ au}}$ describes how many dependent draws correspond to one independent sample



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- new R-hat paper S=NM (in BDA3 N=nm and $n_{E_{\max}}=N/ au$)



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
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 - Warm-up
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- \bullet τ describes how many dependent draws correspond to one independent sample
- new R-hat paper S=NM (in BDA3 N=nm and $n_{E_{\max}}=N/ au$)
- BDA3 focuses on $S_{E_{\max}}$ and not the Monte Carlo error directly new R-hat paper discusses more about MCSEs for

different quantities



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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
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- Diagnostics
 - Warm-up
 - Convegernce
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Estimation of the autocorrelation using several chains

$$\hat{\rho}_n = 1 - \frac{W - \frac{1}{M} \sum_{m=1}^{M} \hat{\rho}_{n,m}}{2\hat{\text{var}}^+}$$



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 - Gibbs sampling
 Metropolis-Hastings
- Diagnostics
 Warm-up
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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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 The above equation is used in Stan 2.18+



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

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 - takes into account if the chains are not mixing (the chains have not converged)
- BDA3 has slightly different and less accurate equation.
 The above equation is used in Stan 2.18+
- Compared to a method which computes the autocorrelation from a single chain, the multi-chain estimate has smaller variance



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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Time series analysis

• Estimation of τ

$$\tau = 1 + 2\sum_{t=1}^{\infty} \hat{\rho}_t$$



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 - Gibbs sampling
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- Diagnostics
 - Warm-up
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- ullet empirical autocorrelation function is noisy and thus estimate of au is noisy
- noise is larger for longer lags (less observations)



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- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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$$\hat{\tau} = 1 + 2\sum_{t=1}^{T} \hat{\rho}_t$$



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$$\hat{\tau} = 1 + 2\sum_{t=1}^{T} \hat{\rho}_t$$

- As τ is estimated from a finite number of draws, it's expectation is overoptimistic
 - if $\hat{\tau} > MN/20$ then the estimate is unreliable



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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
 - S_{eff}, MCSE, and autocorrelation

Geyer's adaptive window estimator

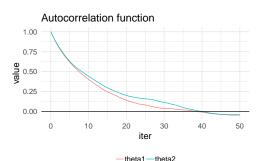
- Truncation can be decided adaptively
 - for stationary, irreducible, recurrent Markov chain
 - let $\Gamma_m = \rho_{2m} + \rho_{2m+1}$, which is sum of two consequent autocorrelations
 - Γ_m is positive, decreasing and convex function of m



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 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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 - let $\Gamma_m = \rho_{2m} + \rho_{2m+1}$, which is sum of two consequent autocorrelations
 - Γ_m is positive, decreasing and convex function of m
- Initial positive sequence estimator (Geyer's IPSE)
 - Choose the largest m so, that all values of the sequence $\hat{\Gamma}_1, \ldots, \hat{\Gamma}_m$ are positive





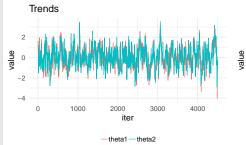
- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
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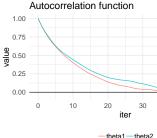
Effective sample size $\mathrm{ESS} = S_{E_{\mathrm{max}}} \approx S/\hat{\tau}$

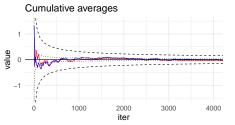


- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
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- Diagnostics
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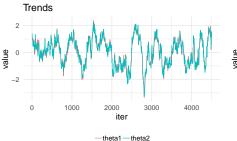


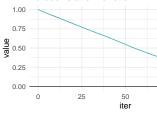
$$\hat{\tau} = 1 + 2 \sum_{t=1}^{T} \hat{\rho}$$
 ≈ 24



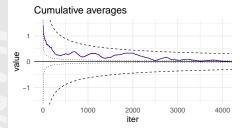
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Autocorrelation function



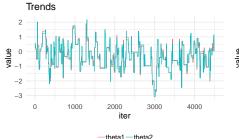
$$\hat{ au} = 1 + 2 \sum_{t=1}^{T} \hat{
ho}$$
 $pprox 104$

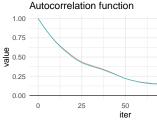
theta1 - theta2



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- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
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Effective sample size $ESS = S_{E_{max}} \approx S/\hat{\tau}$





tneta1 —tneta2

Cumulative averages

1
0
1000 2000 3000 4000
iter

$$\hat{ au} = 1 + 2 \sum_{t=1}^{T} \hat{
ho}_t$$
 $pprox 63$

theta1 - theta2

-theta1 —theta2 - - 95% interval for MCMC error · · · 95% interval for indepen



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 - Gibbs sampling
 - Metropolis-Hastings
- Diagnostics
 - Warm-up
 - Convegernce
 - Convegernce
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- Nonlinear dependencies
 - optimal proposal depends on location



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- Nonlinear dependencies
 - optimal proposal depends on location
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 - optimal proposal depends on location
- Multimodal
 - difficult to move from one mode to another
- Long-tailed with non-finite variance and mean
 - central limit theorem for expectations does not hold