

- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings

Bayesian Statistics and Data Analysis Lecture 5

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$$\begin{split} E_{p(\theta|y)}[f(\theta)] &= \int f(\theta) p(\theta|y) d\theta, \\ \text{where} \quad p(\theta|y) &= \frac{p(y|\theta) p(\theta)}{\int p(y|\theta) p(\theta) d\theta} \end{split}$$



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



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• 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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- Monte Carlo methods we have discussed so far
 - Inverse CDF works for 1D



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- What to do in high dimensions?
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 - Laplace, Variational*, EP* (Ch 4, 13*, next course)



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



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



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



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 - + central limit theorem holds for expectations
 - draws are dependent
 - construction of efficient Markov chains is not always easy



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• 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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- Transition distribution $T_t(\theta_t|\theta_{t-1})$ (may depend on t)



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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(θ|y)



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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 $\theta_{j,t-1} = (\theta_{1,J},\ldots,\theta_{j-1,t},\theta_{j+1,t-1},\ldots,\theta_{t-1,J})$

- *j* can be multiple (blocked) parameters
- 1D sampling (|j| = 1) is generally easy



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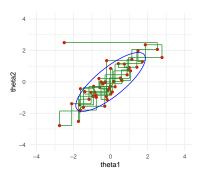
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- *j* can be multiple (blocked) parameters
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- Related to the (stochastic) EM algorithm



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Draws — Steps of the sampler — 90% HPD

demo



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



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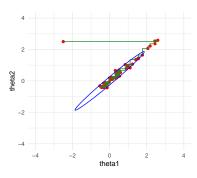
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- No algorithm parameters to tune
- For not so easy conditionals, use e.g. inverse-CDF
- Several parameters can be updated in blocks (blocking)
- Slow if parameters are highly dependent in the posterior...



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Sampling conditional vs joint

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



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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?
 http://elevanth.org/blog/2017/11/28/
 build-a-better-markov-chain/



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- Algorithm
 - 1. starting point θ^0
 - 2. t = 1, 2, ...
 - (a) pick a proposal θ^* from the 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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$$r = \frac{p(\theta^*|y)}{p(\theta^{t-1}|y)}$$



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

$$\theta^t = \begin{cases} \theta^* & \text{with probability min}(r,1) \\ \theta^{t-1} & \text{otherwise} \end{cases}$$



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

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

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



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



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- rejection of a proposal increments the time t also by one
 ie, the new state is the same as previous
- 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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- Example: one bivariate observation (y_1, y_2)
 - bivariate normal distribution with unknown mean and known covariance

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

- proposal distribution $J_t(\theta^*|\theta^{t-1}) = \mathcal{N}(\theta^*|\theta^{t-1}, \sigma_p^2)$
- Demo http://elevanth.org/blog/2017/11/28/ build-a-better-markov-chain/



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 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



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- 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
 - 1. Prove that simulated series is a Markov chain which has unique stationary distribution
 - Prove that this stationary distribution is the desired target distribution



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

c) recurrent / not transient



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- Prove that simulated series is a Markov chain which has unique stationary distribution
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 - positive probability of eventually reaching any state from any other state
 - b) aperiodic
 - = 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
 - holds for a random walk on any proper distribution (except for trivial exceptions)



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



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 - consider any two such points θ_a and θ_b drawn from $p(\theta|y)$ and labeled so that $p(\theta_b|y) \ge p(\theta_a|y)$



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 - consider any two such points θ_a and θ_b drawn from $p(\theta|y)$ and labeled so that $p(\theta_b|y) \ge p(\theta_a|y)$
 - the unconditional probability density of a transition from θ_a to θ_b is

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



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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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 - and labeled so that $p(\theta_b|y) \ge p(\theta_a|y)$
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$$p(\theta^{t} = \theta_{a}, \theta^{t-1} = \theta_{b}) = p(\theta_{b}|y)J_{t}(\theta_{a}|\theta_{b})\left(\frac{p(\theta_{a}|y)}{p(\theta_{b}|y)}\right)$$
$$= p(\theta_{a}|y)J_{t}(\theta_{a}|\theta_{b}),$$

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



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- 2. Prove that this stationary distribution is the desired target distribution $p(\theta|y)$
 - consider starting algorithm at time t-1 with a draw $heta^{t-1} \sim p(heta|y)$
 - consider any two such points θ_a and θ_b drawn from $p(\theta|y)$ and labeled so that $p(\theta_b|y) \geq p(\theta_a|y)$
 - the unconditional probability density of a transition from θ_a to θ_b is

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

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

$$p(\theta^{t} = \theta_{a}, \theta^{t-1} = \theta_{b}) = p(\theta_{b}|y)J_{t}(\theta_{a}|\theta_{b})\left(\frac{p(\theta_{a}|y)}{p(\theta_{b}|y)}\right)$$
$$= p(\theta_{a}|y)J_{t}(\theta_{a}|\theta_{b}),$$

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

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

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



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- 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



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- After the shape has been selected, it is important to select the scale
 - small scale
 - \rightarrow many steps accepted, but the chain moves slowly due to small steps
 - big scale
 - \rightarrow long steps proposed, but many of those rejected and again chain moves slowly



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 - small scale
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 - big scale
 - \rightarrow long steps proposed, but many of those rejected and again chain moves slowly
- Generic rule for rejection rate is 60-90% (but depends on dimensionality and a specific algorithm variation)



- Monte Carlo recap
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Gibbs sampling

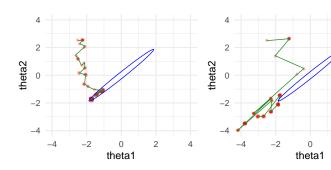
- Specific case of Metropolis-Hastings algorithm
 - single updated (or blocked)
 - proposal distribution is the conditional distribution
 - \rightarrow proposal and target distributions are same
 - ightarrow acceptance probability is 1



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
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 - Metropolis-Hastings

Metropolis

- Usually doesn't scale well to high dimensions
 - if the shape doesn't match the whole distribution, the efficiency drops
 - demo11_2



Draws—Steps of the sampler—90% HPI

Draws—Steps of the sar



- Monte Carlo recap
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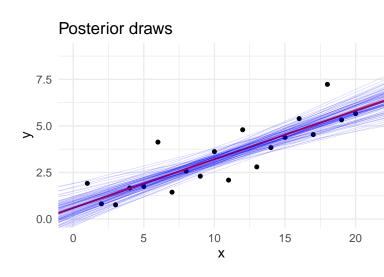
Dynamic Hamiltonian Monte Carlo and NUTS

- Chapter 12 presents some more advanced methods
 - Chapter 12 includes Hamiltonian Monte Carlo and NUTS, which is one of the most efficient methods
 - uses gradient information
 - Hamiltonian dynamic simulation reduces random walk
 - Demo http://elevanth.org/blog/2017/11/28/ build-a-better-markov-chain/



- Monte Carlo recap
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Example of uncertainty in modeling



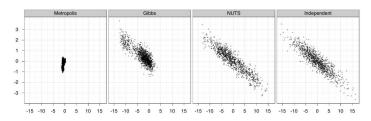


- Monte Carlo recap
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HMC / NUTS

Comparison of algorithms on **highly correlated** 250-dimensional Gaussian distribution

- Do 1,000,000 draws with both Random Walk Metropolis and Gibbs, thinning by 1000
- •Do 1,000 draws using Stan's NUTS algorithm (no thinning)
- •Do 1,000 independent draws (we can do this for multivariate normal)



Source: Jonah Gabry



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings

Warm-up and convergence diagnostics

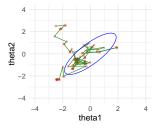
• Asymptotically chain spends the $\alpha\%$ of time where $\alpha\%$ posterior mass is



- Monte Carlo recap
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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



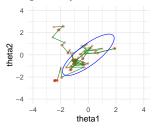
Draws—Steps of the sampler—90% HP



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Warm-up and convergence diagnostics

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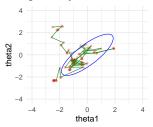
- Draws—Steps of the sampler—90% HPI
- Warm-up = remove draws from the beginning of the chain
 - warm-up may include also phase for adapting algorithm parameters



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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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MCMC draws are dependent

Monte Carlo estimates still valid (central limit theorem holds)

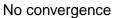
$$E_{p(\theta|y)}[f(\theta)] \approx \frac{1}{S} \sum_{s=1}^{S} f(\theta^{(s)})$$

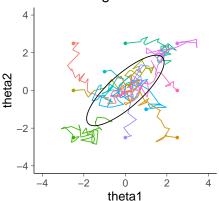
- Estimation of Monte Carlo error is more difficult
 - evaluation of effective sample size



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

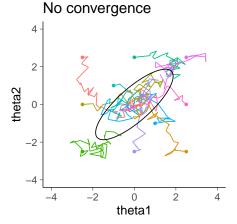






- Monte Carlo recap
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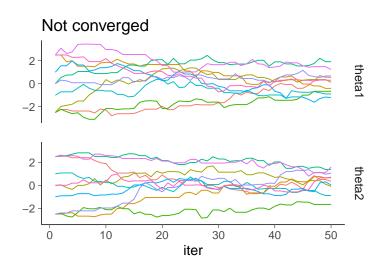
- Use of several chains make convergence diagnostics easier
- Start chains from different starting points preferably overdispersed



 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

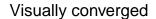


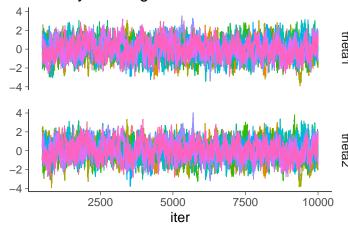
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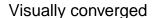
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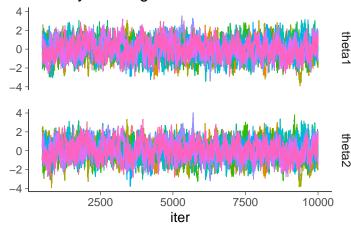






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Visual convergence check is not sufficient



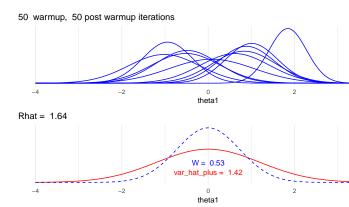
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- BDA3: \hat{R} aka potential scale reduction factor (PSRF)
- Compare means and variances of the chains



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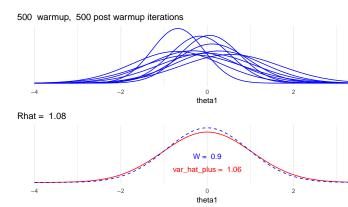
- BDA3: \hat{R} aka potential scale reduction factor (PSRF)
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 W = within chain variance estimate
 var_hat_plus = total variance estimate





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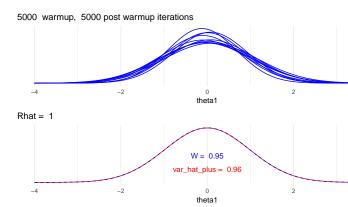
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ullet M chains, each having N draws (with new R-hat notation)



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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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Between chains variance B

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



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- M chains, each having N draws (with new R-hat notation)
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• B/N is variance of the means of the chains



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Between chains variance B

$$\begin{split} B &= \frac{N}{M-1} \sum_{m=1}^{M} (\bar{\theta}_{.m} - \bar{\theta}_{..})^2, \\ \text{where } \bar{\theta}_{.m} &= \frac{1}{N} \sum_{n=1}^{N} \theta_{nm}, \ \bar{\theta}_{..} = \frac{1}{M} \sum_{n=1}^{M} \bar{\theta}_{.m} \end{split}$$

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

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

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

- 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 $N \to \infty$, $E(W) \to var(\theta|y)$
- As $\widehat{\mathrm{var}}^+(\theta|y)$ overestimates and W underestimates, compute

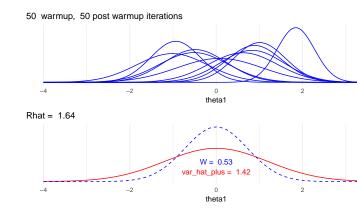
$$\widehat{R} = \sqrt{\frac{\widehat{\text{var}}^+}{W}}$$



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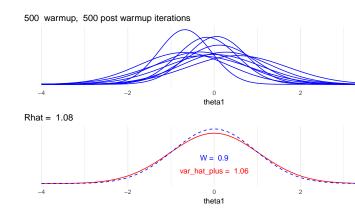




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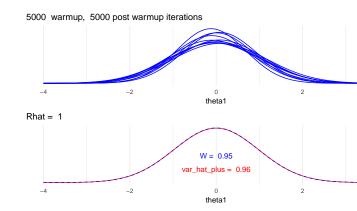




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$$\widehat{R} = \sqrt{\frac{\widehat{\mathrm{var}}^+}{W}}$$

- 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



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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



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Split- \widehat{R}

- BDA3: split- \hat{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



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Rank normalized \widehat{R}

• 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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- 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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- Notation updated compared to BDA3

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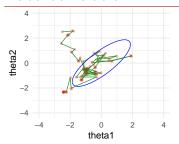
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Time series analysis

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



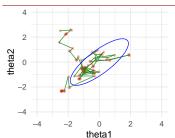
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Draws—Steps of the sampler—90% HPI

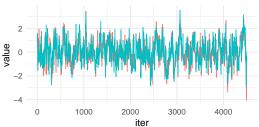


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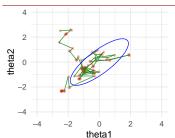
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Trends



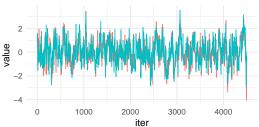


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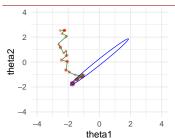
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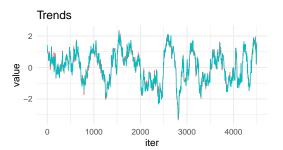




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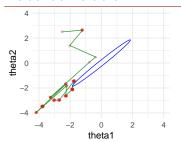
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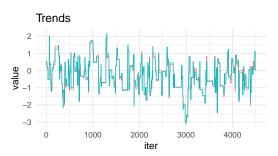


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Auto correlation



• Draws - Steps of the sampler - 90% HPI





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



- Monte Carlo recap
- Markov Chain Monte Carlo (MCMC)
 - Gibbs sampling
 - Metropolis-Hastings

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

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 \bullet τ describes how many dependent draws correspond to one independent sample



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- ullet BDA3 focuses on $S_{E_{
 m max}}$ and not the Monte Carlo error directly new R-hat paper discusses more about MCSEs for

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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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 The above equation is used in Stan 2.18+



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 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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• Estimation of τ

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



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Time series analysis

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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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Geyer's adaptive window estimator

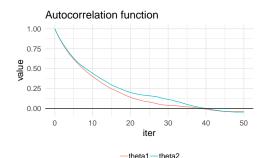
- Truncation can be decided adaptively
 - for stationary, irreducible, recurrent Markov chain
 - let $\Gamma_m = \rho_{2m} + \rho_{2m+1}$, which is sum of two consequent autocorrelations
 - Γ_m is positive, decreasing and convex function of m



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- Initial positive sequence estimator (Geyer's IPSE)
 - Choose the largest m so, that all values of the sequence $\hat{\Gamma}_1, \dots, \hat{\Gamma}_m$ are positive





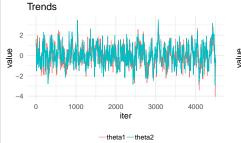
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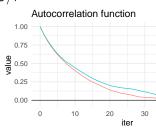
Effective sample size $\mathrm{ESS} = \mathcal{S}_{\mathcal{E}_{\mathrm{max}}} \approx \mathcal{S}/\hat{\tau}$



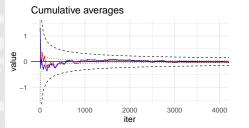
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a1—theta2 —theta1—theta2



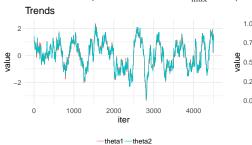
$$\hat{\tau} = 1 + 2 \sum_{t=1}^{T} \hat{\rho}$$
 ≈ 24

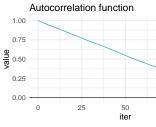
-theta1 —theta2 - - 95% interval for MCMC error -- 95% interval for indepen



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Effective sample size $ESS = S_{E_{max}} \approx S/\hat{\tau}$





Cumulative averages /alue 1000 2000 3000 4000 iter

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

 ≈ 104

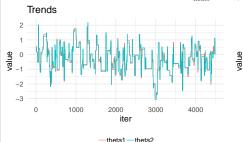
theta1 - theta2

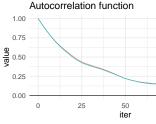
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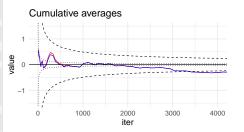
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Effective sample size $ESS = S_{E_{max}} \approx S/\hat{\tau}$





—theta2 —theta1—theta2



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

$$\approx 63$$

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



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- Long-tailed with non-finite variance and mean
 - central limit theorem for expectations does not hold