Chapter 11

- 11.1 Gibbs sampler
- 11.2 Metropolis and Metropolis-Hastings
- 11.3 Using Gibbs and Metropolis as building blocks
- 11.4 Inference and assessing convergence (important)
 - potential scale reduction \hat{R} (R-hat)
- 11.5 Effective number of simulation draws (important)
 - effective sample size (ESS / S_{eff})
- 11.6 Example: hierarchical normal model (quick glance)

Chapter 11 demos

- demo11_1: Gibbs sampling
- demo11_2: Metropolis sampling
- demo11_3: Convergence of Markov chain
- demo11_4: split- \widehat{R} and effective sample size (ESS or $S_{\rm eff}$)

$$E_{p(\theta|y)}[f(\theta)] = \int f(\theta)p(\theta \mid y)d\theta,$$
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• Monte Carlo methods which can sample from $p(\theta^{(s)} \mid y)$ using only $q(\theta^{(s)} \mid y)$

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

Monte Carlo

- Monte Carlo methods we have discussed so far
 - Inverse CDF works for 1D
 - Analytic transformations work for only certain distributions
 - Factorization works only for certain joint distributions
 - Grid evaluation and sampling works in a few dimensions
 - Rejection sampling works mostly in 1D (truncation is a special case)
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 - Importance sampling is reliable only in special cases
- What to do in high dimensions?
 - Markov chain Monte Carlo (Ch 11-12)
 - Laplace, Variational*, EP* (Ch 4,13*)

- Andrey Markov proved weak law of large numbers and central limit theorem for certain dependent-random sequences, which were later named Markov chains
 - CLT saying the sum / mean converges towards normal if the variance is finite (we come back to this in the end of the course)

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 - Deep learning language models are super big Markov models

• Example of a simple Markov chain

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 - draws are dependent
 - construction of efficient Markov chains is not always easy

• Set of random variables $\theta^1, \theta^2, \ldots$, so that with all values of t, θ^t depends only on the previous $\theta^{(t-1)}$

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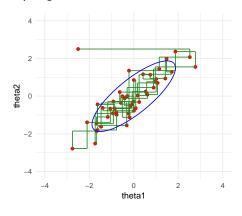
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- Chain has to be initialized with some starting point θ^0
- Transition distribution $T_t(\theta^t \mid \theta^{t-1})$ (may depend on t)
 - by choosing a suitable transition distribution, the stationary distribution of Markov chain is $p(\theta \mid y)$

- Alternate sampling from 1D conditional distributions
 - e.g. normal distribution, sample alternating from $p(\mu \mid \sigma^2, y)$ and $p(\sigma^2 \mid \mu, y)$

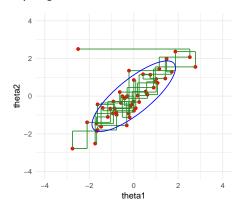
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 - 1D is easy even if no conjugate prior and analytic posterior

- Alternate sampling from 1D conditional distributions
- demo11_1



Draws — Steps of the sampler — 90% HPD

- Alternate sampling from 1D conditional distributions
- demo11 1



Draws — Steps of the sampler — 90% HPD

Basic algorithm

sample
$$\theta_j^t$$
 from $p(\theta_j \mid \theta_{-j}^{t-1}, y)$,
where $\theta_{-j}^{t-1} = (\theta_1^t, \dots, \theta_{j-1}^t, \theta_{j+1}^{t-1}, \dots, \theta_d^{t-1})$

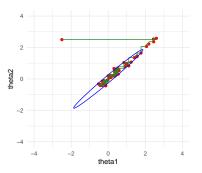
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- Slow if parameters are highly dependent in the posterior
 - demo11₋1 continues



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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?

Metropolis algorithm

- Algorithm
 - 1. starting point θ^0
 - 2. t = 1, 2, ...
 - (a) pick a proposal θ^* from the proposal distribution $J_t(\theta^* \mid \theta^{t-1})$. Proposal distribution has to be symmetric, i.e.

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

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

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- $p(\theta^* \mid y)$ and $p(\theta^{t-1} \mid y)$ have the same normalization terms, and thus instead of $p(\cdot \mid y)$, unnormalized $q(\cdot \mid y)$ can be used, as the normalization terms cancel out!

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

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

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- More examples https://chi-feng.github.io/mcmc-demo/

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Theoretically

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

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

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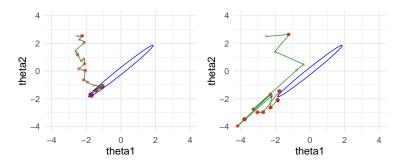
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- Generic rule for rejection rate is 60-90% (but depends on dimensionality and a specific algorithm variation)

Gibbs sampling

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

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

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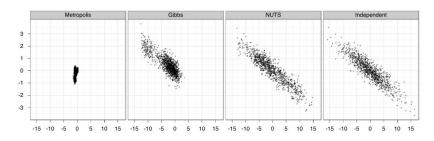
Dynamic Hamiltonian Monte Carlo and NUTS

- 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
 - state-of-the-art MCMC used by most modern probabilistic programming frameworks
- More next week

HMC / NUTS

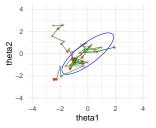
Comparison of algorithms on **highly correlated** 250-dimensional Gaussian distribution

- •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)



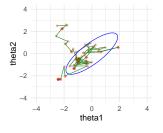
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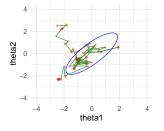
Draws—Steps of the sampler—90% HP

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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
 - Is the sample representative of the target distribution?

MCMC draws are dependent

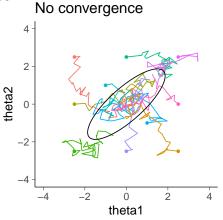
Monte Carlo estimates still valid (central limit theorem holds)

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

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

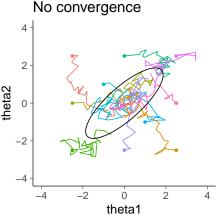
Several chains

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



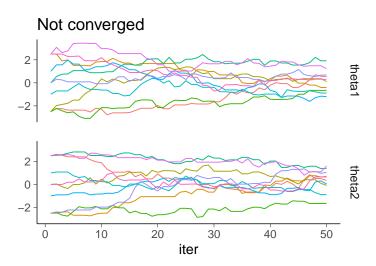
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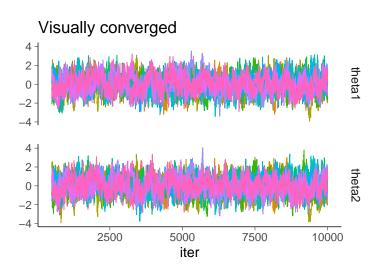


 Remove draws from the beginning of the chains and run chains long enough so that it is not possible to distinguish where each chain started and the chains are well mixed

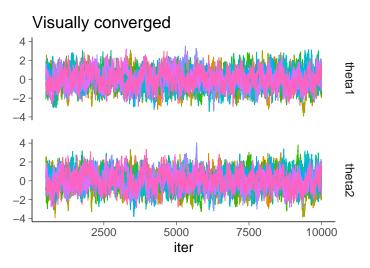
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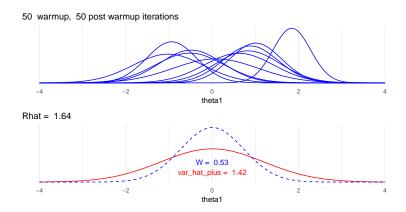
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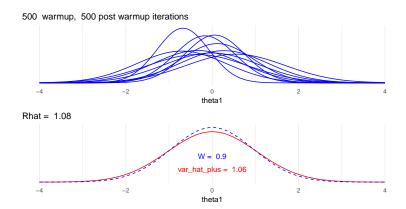
Visual convergence check is not sufficient

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

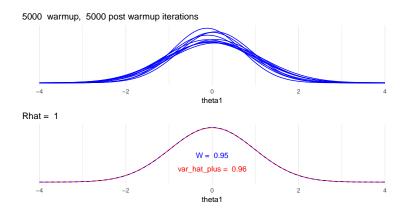
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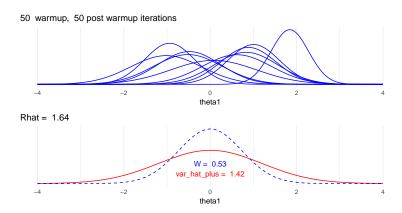
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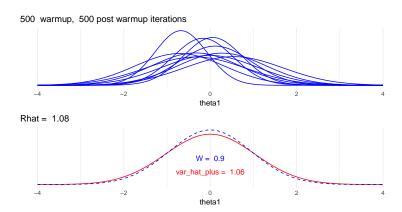
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 - single chains have not yet visited all points in the distribution
 - when $N \to \infty$, $E(W) \to var(\theta \mid y)$
- As $\widehat{\text{var}}^+(\theta \mid y)$ overestimates and W underestimates, compute

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

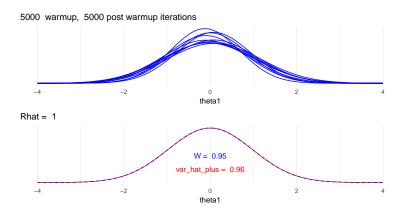
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- if \widehat{R} is big (e.g., R > 1.01), keep sampling
- If R close to 1, it is still possible that chains have not converged
 - if starting points were not overdispersed
 - distribution far from normal (especially if infinite variance)
 - just by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

Split- \widehat{R}

- BDA3: split-R̂
- 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

• 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 \widehat{R} for assessing convergence of MCMC. Bayesian Analysis, doi:10.1214/20-BA1221. https://projecteuclid.org/euclid.ba/1593828229.

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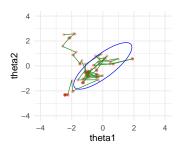
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- Notation updated compared to BDA3

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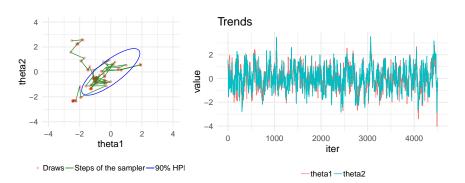
- Autocorrelation function
 - describes the correlation given a certain lag
 - can be used to compare efficiency of MCMC algorithms and parameterizations

Autocorrelation

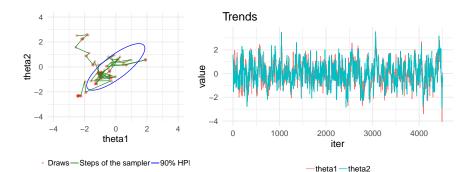


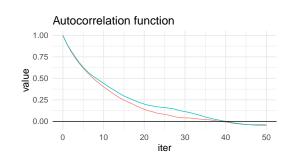
Draws—Steps of the sampler—90% HPI

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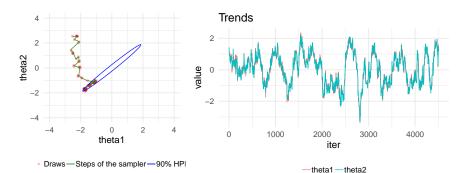


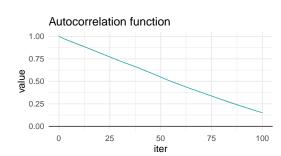
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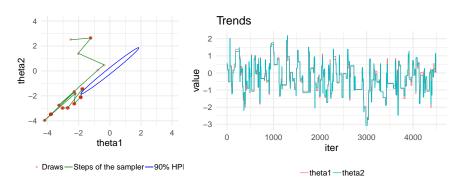


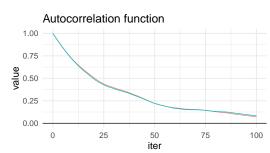
Autocorrelation (slow mixing due to small step size)





Autocorrelation (slow mixing due to many rejections)

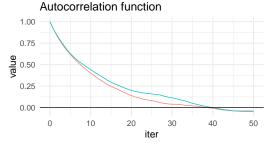




- Time series analysis can be used to estimate Monte Carlo error in case of MCMC
- For expectation $\bar{\theta}$

$$\mathsf{Var}[\bar{ heta}] = rac{\sigma_{ heta}^2}{\mathsf{S}_{\mathsf{eff}}}$$

where $S_{\rm eff} = S/\tau$ (=ESS), and τ is sum of autocorrelations



theta1 — theta2

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- new \widehat{R} paper S = NM (in BDA3 N = nm and $n_{\text{eff}} = N/\tau$)
- BDA3 focuses on $S_{\rm eff}$ and not the Monte Carlo error directly new \widehat{R} paper discusses more about MCSEs for different quantities

Estimation of the autocorrelation using several chains

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

where $\hat{\rho}_{n,m}$ is autocorrelation at lag n for chain m

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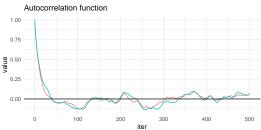
- This combines \widehat{R} and autocorrelation estimates
 - takes into account if the chains are not mixing (the chains have not converged)
- BDA3 has slightly different and less accurate equation. The above equation is used in Stan 2.18+
- Compared to a method which computes the autocorrelation from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

• Estimation of τ

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

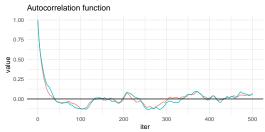
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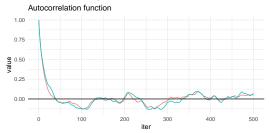
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- less noisy estimate is obtained by truncating

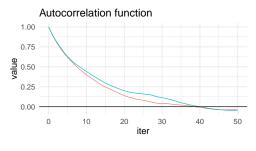
$$\hat{\tau} = 1 + 2\sum_{t=1}^{T} \hat{\rho}_t$$

Geyer's adaptive window estimator

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

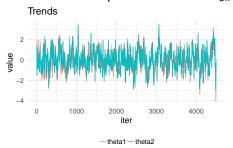
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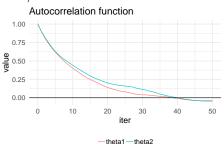
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 - let $\Gamma_m = \rho_{2m} + \rho_{2m+1}$, which is sum of two consequent autocorrelations
 - Γ_m is positive, decreasing and convex function of m
- Initial positive sequence estimator (Geyer's IPSE)
 - Choose the largest m so, that all values of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

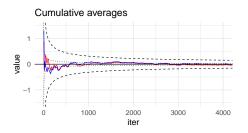


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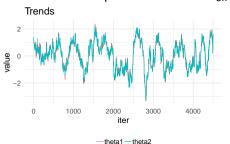


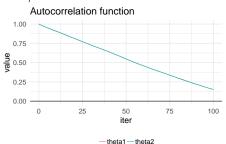


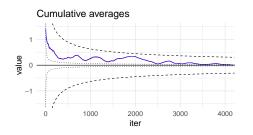


$$\hat{\tau} = 1 + 2 \sum_{t=1}^{T} \hat{\rho}_t$$
 ≈ 24

Effective sample size ESS = $S_{\rm eff} \approx S/\hat{\tau}$



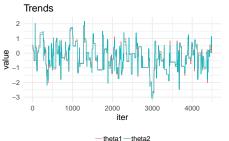


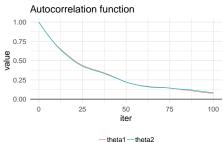


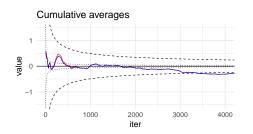
$$\hat{\tau} = 1 + 2\sum_{t=1}^{T} \hat{\rho}$$

$$\approx 104$$

Effective sample size ESS = $S_{\rm eff} \approx S/\hat{ au}$







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

$$\approx 63$$

- Nonlinear dependencies
 - · optimal proposal depends on location

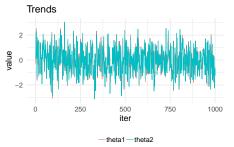
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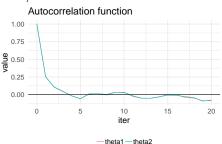
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- Long-tailed with non-finite variance and mean
 - central limit theorem for expectations does not hold

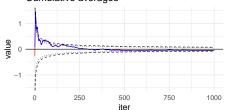
Next week: HMC, NUTS, and dynamic HMC

Effective sample size ESS = $S_{\rm eff} \approx S/\hat{\tau}$





Cumulative averages



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

$$\approx 1.6$$

Further diagnostics

- Dynamic HMC/NUTS has additional diagnostics
 - divergences
 - tree depth exceedences
 - fraction of missing information