#### 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  $\hat{R}$
- 11.5 Effective number of simulation draws (important)
  - effective sample size N<sub>eff</sub>
- 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  $N_{\rm eff}$

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

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

#### 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 less than a few dimensions
  - Rejection sampling works mostly in 1D (truncation is a special case)
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  - Rejection sampling works mostly in 1D (truncation is a special case)
  - 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)

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- Markov's one example was the sequence of letters in Pushkin's novel "Yevgeniy Onegin"

• Example of a simple Markov chain on black board

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

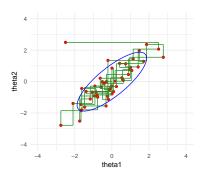
• Set of random variables  $\theta^1, \theta^2, \ldots$ , so that with all values of t,  $\theta^t$  depends only on the previous  $\theta^{(t-1)}$ 

$$p(\theta^t|\theta^1,\ldots,\theta^{(t-1)})=p(\theta^t|\theta^{(t-1)})$$

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

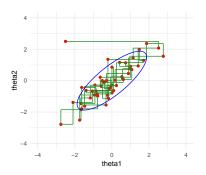
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Draws — Steps of the sampler — 90% HPD

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- demo11 1



- Draws Steps of the sampler 90% HPD
- Basic algorithm

sample 
$$\theta_j^t$$
 from  $p(\theta_j | \theta_{-j}^{t-1}, y)$ ,  
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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  - BUGS/WinBUGS/OpenBUGS/JAGS

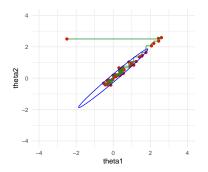
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- Slow if parameters are highly dependent in the posterior

demo11\_1 continues



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- Can we use that to form a Markov chain? http://elevanth.org/blog/2017/11/28/build-a-better-markov-chain/

- Algorithm
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  - 2. t = 1, 2, ...
    - (a) pick a proposal  $\theta^*$  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  $\theta_a$ ,  $\theta_b$

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

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- instead of  $p(\theta|y)$ , unnormalized  $q(\theta|y)$  can be used, as the normalization terms cancel out!

## Metropolis algorithm

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

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

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

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#### 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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    - = aperiodic (return times are not periodic)
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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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$$\begin{aligned}
\rho(\theta^t = \theta_a, \theta^{t-1} = \theta_b) &= \rho(\theta_b | y) J_t(\theta_a | \theta_b) \left( \frac{\rho(\theta_a | y)}{\rho(\theta_b | y)} \right) \\
&= \rho(\theta_a | y) J_t(\theta_a | \theta_b),
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which is the same as the probability of transition from  $\theta_a$  to  $\theta_b$ , since we have required that  $J_t(\cdot|\cdot)$  is symmetric

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

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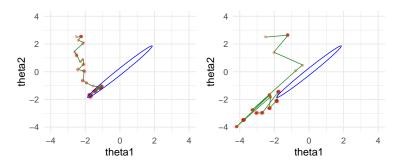
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- Generic rule for rejection rate is 60-90% (but depends on dimensionality and a specific algorithm variation)

# 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

- Ususally 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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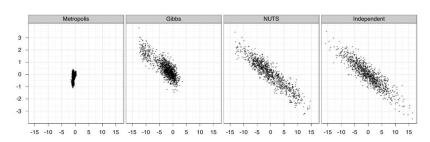
#### Dynamic Hamiltonian Monte Carlo and NUTS

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

#### HMC / NUTS

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

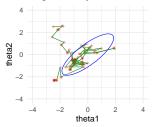
- •Do **1,000,000** draws with both Random Walk Metropolis and Gibbs, thinning by 1000
- •Do 1,000 draws using Stan's NUTS algorithm (no thinning)
- •Do 1,000 independent draws (we can do this for multivariate normal)



Source: Jonah Gabry

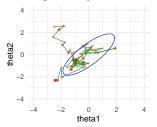
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  - but in finite time the initial part of the chain may be non-representative and lower error of the estimate can be obtained by throwing it away



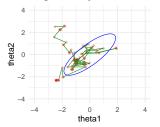
Draws—Steps of the sampler—90% HP

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  - 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



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  - warm-up may include also phase for adapting algorithm parameters

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- 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?

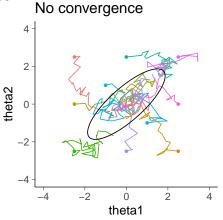
#### MCMC draws are dependent

Monte Carlo estimates still valid (central limit theorem holds)

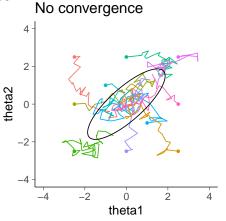
$$E_{p(\theta|y)}[f(\theta)] \approx \frac{1}{S} \sum_{s=1}^{S} f(\theta^{(s)})$$

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

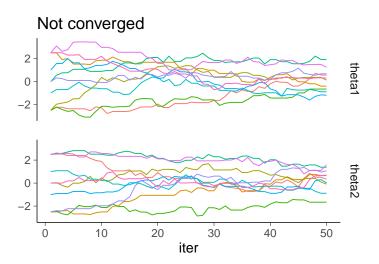
- Use of several chains make convergence diagnostics easier
- Start chains from different starting points preferably overdispersed

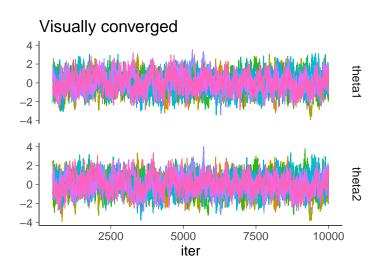


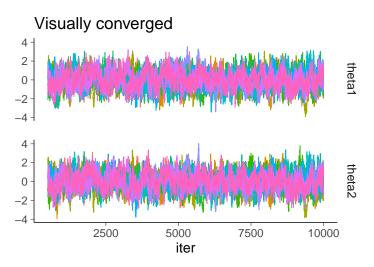
- Use of several chains make convergence diagnostics easier
- Start chains from different starting points preferably overdispersed



 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







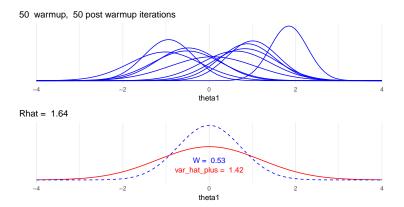
Visual convergence check is not sufficient

# $\widehat{R}$ : comparison of within and between variances of the chains

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

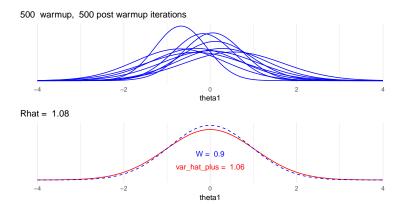
# $\widehat{R}$ : comparison of within and between variances of the chains

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   var hat plus = total variance estimate



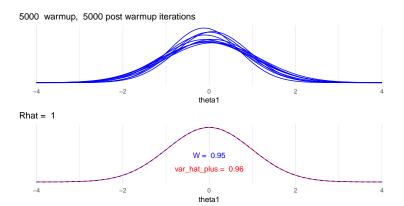
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Within chains variance W

$$W = \frac{1}{m} \sum_{i=1}^{m} s_j^2$$
, where  $s_j^2 = \frac{1}{n-1} \sum_{i=1}^{n} (\psi_{ij} - \bar{\psi}_{.j})^2$ 



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Between chains variance B

$$B = \frac{n}{m-1} \sum_{j=1}^{m} (\bar{\psi}_{.j} - \bar{\psi}_{..})^2, \text{ where } \bar{\psi}_{.j} = \frac{1}{n} \sum_{i=1}^{n} \psi_{ij}, \bar{\psi}_{..} = \frac{1}{m} \sum_{j=1}^{m} \bar{\psi}_{.j}$$

• B/n is variance of the means of the chains



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

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



Estimate total variance var(ψ|y) as a weighted mean of W and B

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- Given finite n, W underestimates marginal posterior variance
  - single chains have not yet visited all points in the distribution
  - when  $n \to \infty$ ,  $E(W) \to var(\psi|y)$



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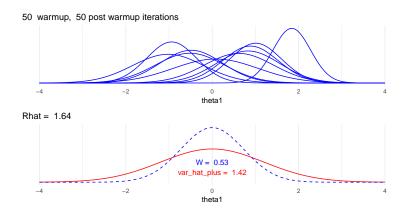
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- Given finite n, W underestimates marginal posterior variance
  - single chains have not yet visited all points in the distribution
  - when  $n \to \infty$ ,  $E(W) \to var(\psi|y)$
- As  $\widehat{\text{var}}^+(\psi|y)$  overestimates and W underestimates,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

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

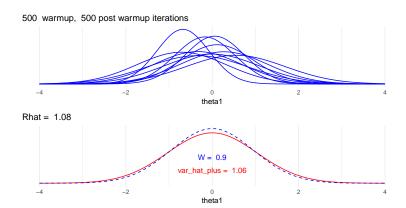


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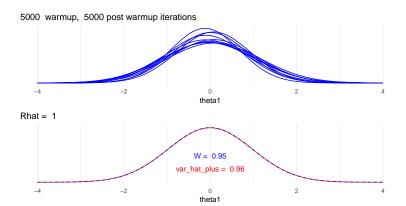


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- Estimates how much the scale of  $\psi$  could reduce if  $n \to \infty$
- $R \to 1$ , when  $n \to \infty$
- if R is big (e.g., R > 1.01), keep sampling



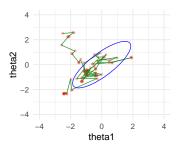
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- $R \to 1$ , when  $n \to \infty$
- if R is big (e.g., R > 1.01), keep sampling
- If R close to 1, it is still possible that chains have not converged
  - if starting points were not overdispersed
  - distribution far from normal (especially if infinite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

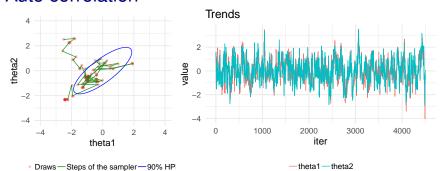
# Split-R

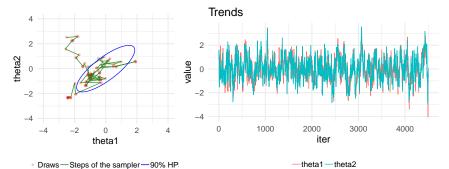
- BDA3: split-R̂
- Examines mixing and stationarity of chains
- To examine stationarity chains are splitted to two parts
  - after splitting, we have *m* chains, each having *n* draws
  - scalar draws  $\psi_{ij}$   $(i = 1, \dots, n; j = 1, \dots, m)$
  - compare means and variances of the split chains

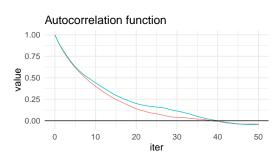
- Auto correlation function
  - describes the correlation given a certain lag
  - can be used to compare efficiency of MCMC algorithms and parameterizations

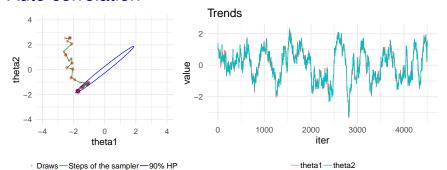


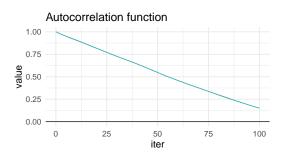
Draws—Steps of the sampler—90% HP

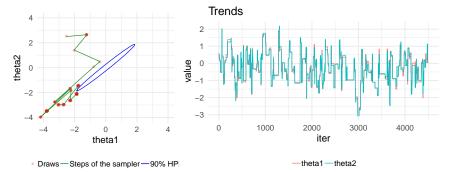


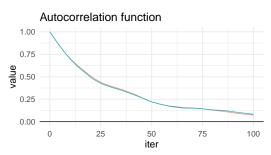












- Time series analysis can be used to estimate Monte Carlo error in case of MCMC
- For expectation  $\bar{\theta}$

$$\operatorname{Var}[\bar{\theta}] = \frac{\sigma_{\theta}^2}{N/\tau}$$

where  $\tau$  is sum of autocorrelations

- $\bullet$   $\tau$  describes how many dependent draws correspond to one independent sample
- in BDA3 N = nm
- $n_{\rm eff} = nm/\tau$
- BDA3 focuses on n<sub>eff</sub> and not the Monte Carlo error directly

Estimation of the autocorrelation using several chains

$$\hat{\rho}_t = 1 - \frac{W - \frac{1}{M} \sum_{j=1}^{m} \hat{\rho}_{t,j}}{2\widehat{\text{var}}^+}$$

where  $\hat{\rho}_{t,j}$  is autocorrelation at lag t for chain j

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 BDA3 has slightly different less accurate equation. The above equation is used in Stan 2.18+

Estimation of the autocorrelation using several chains

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where  $\hat{\rho}_{t,j}$  is autocorrelation at lag t for chain j

- BDA3 has slightly different less accurate equation. The above equation is used in Stan 2.18+
- Compared to usual method which computes the autocorrelation from a single chain, this estimate has smaller variance

• Estimation of  $\tau$ 

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

where  $\hat{\rho}_t$  is empirical autocorrelation

- $\bullet$  empirical autocorrelation function is noisy and thus estimate of  $\tau$  is noisy
- noise is larger for longer lags (less observations)
- less noisy estimate is obtained by truncating

$$au pprox 1 + 2\sum_{t=1}^{T} \hat{
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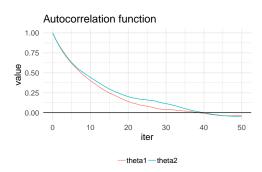
- As  $\tau$  is estimated from a finite number of draws, it's expectation is overoptimistic
  - if  $\tau > mn/20$  then the estimate is unreliable

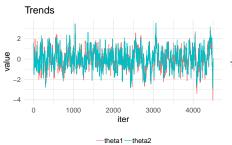
## Geyer's adaptive window estimator

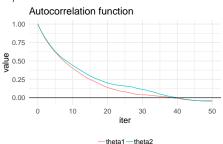
- Truncation can be decided adaptively
  - for stationary, irreducible, recurrent Markov chain
  - let  $\Gamma_m = \rho_{2m} + \rho_{2m+1}$ , which is sum of two consequent autocorrelations
  - $\Gamma_m$  is positive, decreasing and convex function of m

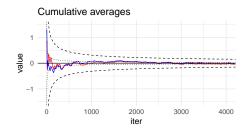
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  - $\Gamma_m$  is positive, decreasing and convex function of m
- Initial positive sequence estimator (Geyer's IPSE)
  - Choose the largest m so, that all values of the sequence  $\hat{\Gamma}_1, \dots, \hat{\Gamma}_m$  are positive

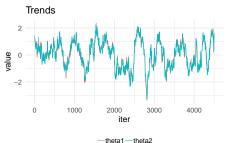


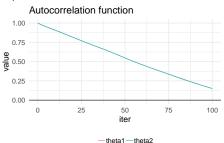


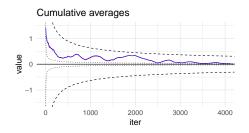




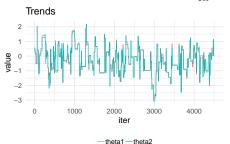
$$au pprox 1 + 2\sum_{t=1}^{T} \hat{
ho}_t$$
  $pprox 24$ 

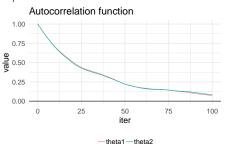


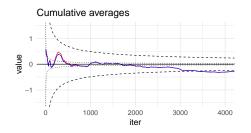




$$\tau \approx 1 + 2 \sum_{t=1}^{T} \hat{\rho}_{t}$$
$$\approx 104$$







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

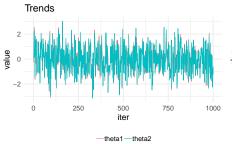
- Nonlinear dependencies
  - optimal proposal depends on location

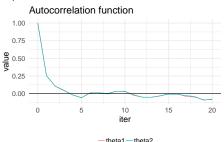
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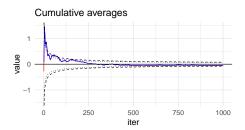
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- Multimodal
  - difficult to move from one mode to another
- Long-tailed with non-finite variance and mean
  - central limit theorem for expectations does not hold

# Next week: HMC, NUTS, and dynamic HMC







$$au pprox 1 + 2 \sum_{t=1}^{T} \hat{
ho}_t$$
  $pprox 1.6$ 

## Further diagnostics

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