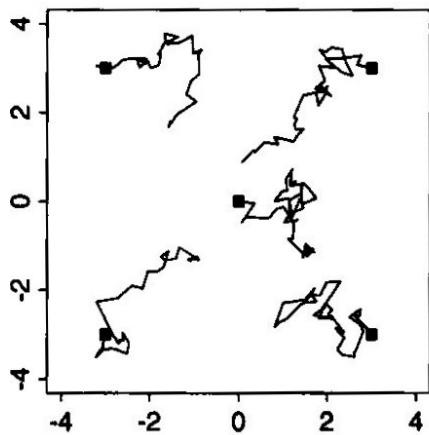


# Markov Chain Monte Carlo

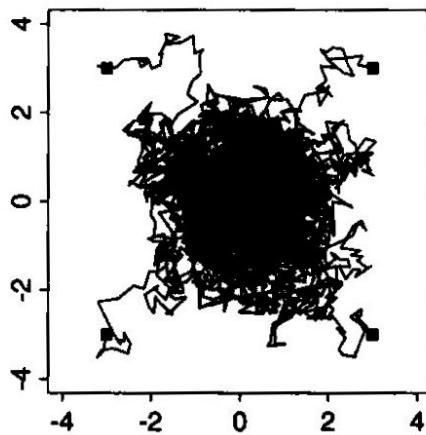
## Simulation for bi-variate standard normal distribution

$$P(\theta^t | \theta^{t-1}) = N(\theta^{t-1}, 0.2^2 I)$$

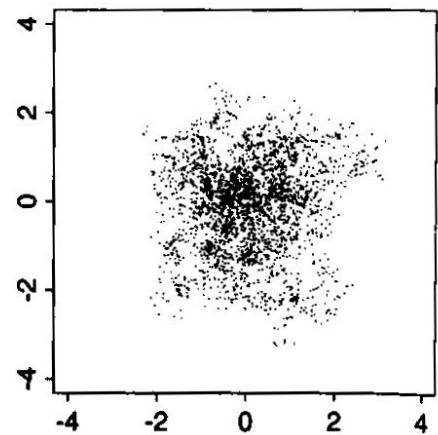
50 iterations



1000 iterations



last 500 samples



Number of  
iterations

$\theta_1$

95% intervals and  $\hat{R}$  for ...

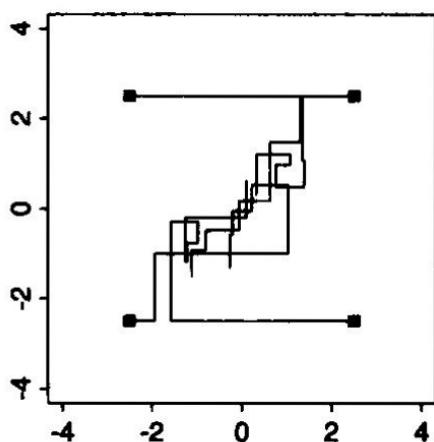
$\theta_2$

$\log p(\theta_1, \theta_2 | y)$

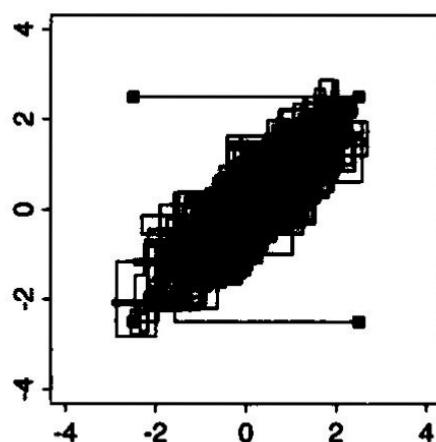
Number of iterations	$\theta_1$	$\theta_2$	$\log p(\theta_1, \theta_2   y)$
50	[-2.14, 3.74], 12.3	[-1.83, 2.70], 6.1	[-8.71, -0.17], 6.1
500	[-3.17, 1.74], 1.3	[-2.17, 2.09], 1.7	[-5.23, -0.07], 1.3
2000	[-1.83, 2.24], 1.2	[-1.74, 2.09], 1.03	[-4.07, -0.03], 1.10
5000	[-2.09, 1.98], 1.02	[-1.90, 1.95], 1.03	[-3.70, -0.03], 1.00
$\infty$	[-1.96, 1.96], 1	[-1.96, 1.96], 1	[-3.69, -0.03], 1

Gibbs for normal distribution  $\rho=0.8$ ,  $(y_1, y_2)=(0,0)$

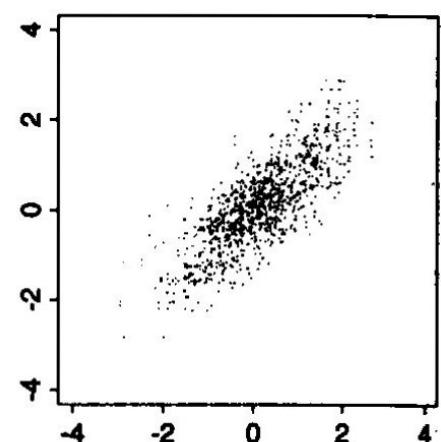
10 iterations



500 iterations



last 250 samples



# Hierarchical Normal Model

Congulation time (seconds) for randomly drawn blood

Diet Measurements

A	62, 60, 63, 59
B	63, 67, 71, 64, 65, 66
C	68, 66, 71, 67, 68, 68
D	56, 62, 60, 61, 63, 64, 63, 59

Posterior from 10 Gibbs sequences of length 100

Estimand	Posterior quantiles					$\widehat{R}$
	2.5%	25%	median	75%	97.5%	
$\theta_1$	58.9	60.6	61.3	62.1	63.5	1.01
$\theta_2$	63.9	65.3	65.9	66.6	67.7	1.01
$\theta_3$	66.0	67.1	67.8	68.5	69.5	1.01
$\theta_4$	59.5	60.6	61.1	61.7	62.8	1.01
$\mu$	56.9	62.2	63.9	65.5	73.4	1.04
$\sigma$	1.8	2.2	2.4	2.6	3.3	1.00
$\tau$	2.1	3.6	4.9	7.6	26.6	1.05
$\log p(\mu, \log \sigma, \log \tau   y)$	-67.6	-64.3	-63.4	-62.6	-62.0	1.02
$\log p(\theta, \mu, \log \sigma, \log \tau   y)$	-70.6	-66.5	-65.1	-64.0	-62.4	1.01