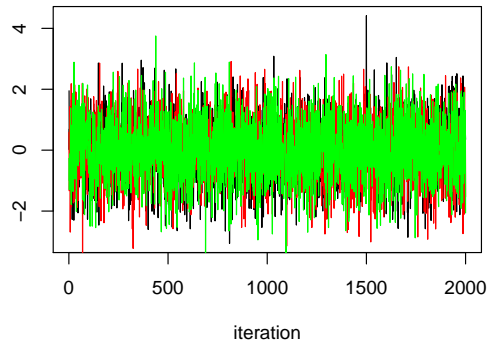
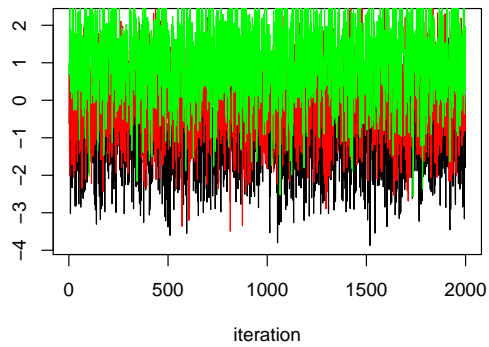


Bayes: Homework 8
Leslie Gains-Germain

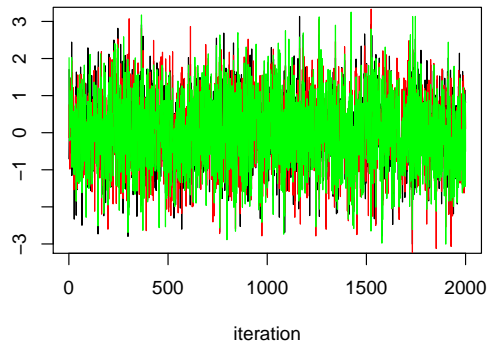
1. (a) i. Scenario 1: three chains from $N(0, 1)$ white noise.



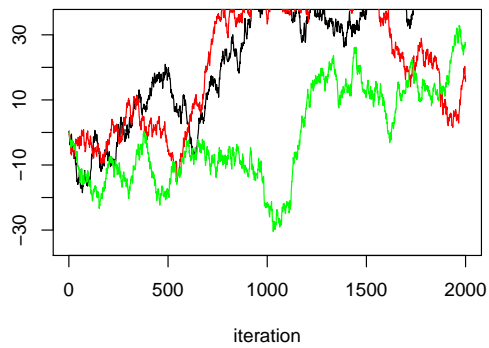
- ii. Scenario 2: one chain from $N(-1, 1)$, one chain from $N(0, 1)$, and one chain from $N(1, 1)$.



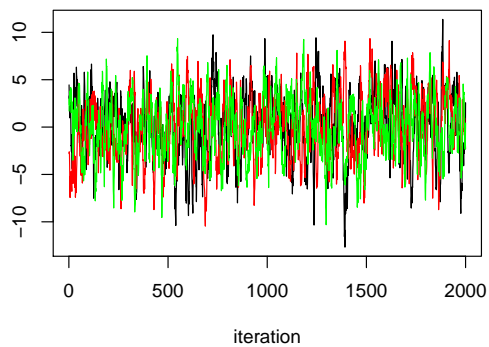
- iii. Scenario 3: three from $MVN(0, \Sigma)$ with common $\sigma^2 = 1$ and $\rho = 0.8$.



iv. Scenario 4: three from non-stationary correlated chains using `diffinv(rnorm(999))`.



v. Scenario 5: three from stationary chains with correlation using
`filter(rnorm(1000), filter=rep(1, 10), circular=TRUE)`.



(b) The effective sample size is calculated differently in the newest version of the `coda` package than described in the textbook. In the textbook, the effective sample size is dividing

the number of iterations (over all chains) by a formula that measures autocorrelation, so that chains with higher autocorrelation have lower effective sample sizes. In the newest version of the `coda` package, however, the number of iterations is divided by the spectral density. Details are on page 286 of the BDA3 text and the Effective Sample size section of the convergence diagnostic document.

In the `coda` package, \hat{R} is calculated in the same way as the described in Chapter 11 in the BDA3 textbook.

- (c) The below table shows the \hat{R} and effective sample sizes for each scenario.

scenario	neff	rhat
1	5674.27	1.00
2	6291.69	1.74
3	5659.49	1.00
4	7.34	2.22
5	540.03	1.00

- (d) The table below shows the z-statistics for all three chains from Geweke's diagnostic. The last three columns show the results from Raftery and Lewis's diagnostic (burn in and iterations required as well as the dependence factor).

scenario	zstats1	zstats2	zstats3	burn.in	n.iter	dependence.factor
1	0.00	-1.87	-0.60	2.00	3660.00	0.98
2	-0.17	-1.17	0.12	3.00	4255.00	1.14
3	-0.26	-0.05	0.33	2.00	3866.00	1.03
4	-11.17	-3.51	-2.35	198.00	215594.00	57.60
5	0.15	-2.19	-0.09	20.00	21264.00	5.68

2.

3. (a) i. My R function to calculate the density of a folded non-central t-distribution is shown below.

```
# function to calculate density of a folded non-central t-distribution
```

ii.