1. (a)

- 주어진 밀도 함수 f(x)와 제안 분포 q(x)~BETA(2,3)를 사용하여 중요도 샘플링을 진행
- 중요도 추출의 기본적인 방법은 제안 분포에서 샘플을 생성한 후, 이 샘플을 이용해 기댓값을 추정하는 것
- 알고리듬
 - 1) $x_i \sim BETA(2,3)$ 에서 샘플 $x_1 x_2 ... x_n$ 을 생성
 - 2) $w_i = \frac{f(x_i)}{g(x_i)}$ 를 계산하여 각 샘플의 중요도 가중치를 구함
 - 3) 기댓값 E[f(x)]는 중요도 가중치로 평균을 계산하여 구함

$$E[f(x)] pprox rac{\displaystyle\sum_{i=1}^{i=1} w_i x_i}{\displaystyle\sum_{i=1}^{i=1} w_i}$$

(b) R로 구현하여 기댓값 구하기

```
# f(x) 정의
f <- function(x) {
+ exp(-x) * x^(2 - 1) * (1 - x)^(3 - 1)
+ }

# 제안 분포 g(x) : BETA(2, 3)에서 샘플링
n <- 10000
x_samples <- rbeta(n, 2, 3)

# 중요도 비율 계산
w <- f(x_samples) / dbeta(x_samples, 2, 3)

# 기댓값 계산
expectation <- sum(w * x_samples) / sum(w)
print(expectation)
[1] 0.363246
```

```
# R 코드에서 필요한 패키지 로드
library(rstan)
# 데이터 준비
data list \leftarrow list(N = 10, x = c(3, 5, 3, 4, 4, 2, 3, 5, 7, 4))
# Stan 모델 컴파일 및 샘플링
fit <- stan(model_code = stan_model_code, data = data_list, iter = 15000, warmup =
5000, chains = 4)
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 1).
Chain 1:
Chain 1: Gradient evaluation took 3.6e-05 seconds
Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.36
seconds.
Chain 1: Adjust your expectations accordingly!
Chain 1:
Chain 1:
Chain 1: Iteration:
                     1 / 15000 [ 0%] (Warmup)
Chain 1: Iteration: 1500 / 15000 [ 10%] (Warmup)
Chain 1: Iteration: 3000 / 15000 [ 20%] (Warmup)
Chain 1: Iteration: 4500 / 15000 [ 30%] (Warmup)
Chain 1: Iteration: 5001 / 15000 [ 33%] (Sampling)
Chain 1: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 1: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 1: Iteration: 9500 / 15000 [ 63%] (Sampling)
Chain 1: Iteration: 11000 / 15000 [ 73%] (Sampling)
Chain 1: Iteration: 12500 / 15000 [ 83%] (Sampling)
Chain 1: Iteration: 14000 / 15000 [ 93%] (Sampling)
Chain 1: Iteration: 15000 / 15000 [100%] (Sampling)
Chain 1:
Chain 1: Elapsed Time: 0.053 seconds (Warm-up)
Chain 1:
                       0.112 seconds (Sampling)
Chain 1:
                       0.165 seconds (Total)
Chain 1:
```

```
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 2).
Chain 2:
Chain 2: Gradient evaluation took 7e-06 seconds
Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.07
seconds.
Chain 2: Adjust your expectations accordingly!
Chain 2:
Chain 2:
Chain 2: Iteration:
                     1 / 15000 [ 0%] (Warmup)
Chain 2: Iteration: 1500 / 15000 [ 10%] (Warmup)
Chain 2: Iteration: 3000 / 15000 [ 20%] (Warmup)
Chain 2: Iteration: 4500 / 15000 [ 30%] (Warmup)
Chain 2: Iteration: 5001 / 15000 [ 33%] (Sampling)
Chain 2: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 2: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 2: Iteration: 9500 / 15000 [ 63%] (Sampling)
Chain 2: Iteration: 11000 / 15000 [ 73%] (Sampling)
Chain 2: Iteration: 12500 / 15000 [ 83%] (Sampling)
Chain 2: Iteration: 14000 / 15000 [ 93%] (Sampling)
Chain 2: Iteration: 15000 / 15000 [100%] (Sampling)
Chain 2:
Chain 2: Elapsed Time: 0.057 seconds (Warm-up)
Chain 2:
                       0.123 seconds (Sampling)
Chain 2:
                       0.18 seconds (Total)
Chain 2.
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 3).
Chain 3:
Chain 3: Gradient evaluation took 4e-06 seconds
Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.04
seconds.
Chain 3: Adjust your expectations accordingly!
Chain 3:
Chain 3:
Chain 3: Iteration:
                     1 / 15000 [ 0%] (Warmup)
```

```
Chain 3: Iteration:
                  1500 / 15000 [ 10%]
                                        (Warmup)
Chain 3: Iteration:
                  3000 / 15000 [ 20%] (Warmup)
Chain 3: Iteration: 4500 / 15000 [ 30%] (Warmup)
Chain 3: Iteration: 5001 / 15000 [ 33%] (Sampling)
Chain 3: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 3: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 3: Iteration: 9500 / 15000 [ 63%] (Sampling)
Chain 3: Iteration: 11000 / 15000 [ 73%] (Sampling)
Chain 3: Iteration: 12500 / 15000 [ 83%] (Sampling)
Chain 3: Iteration: 14000 / 15000 [ 93%] (Sampling)
Chain 3: Iteration: 15000 / 15000 [100%]
                                        (Sampling)
Chain 3:
Chain 3: Elapsed Time: 0.05 seconds (Warm-up)
Chain 3:
                       0.124 seconds (Sampling)
Chain 3:
                        0.174 seconds (Total)
Chain 3:
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 4).
Chain 4:
Chain 4: Gradient evaluation took 4e-06 seconds
Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.04
seconds.
Chain 4: Adjust your expectations accordingly!
Chain 4:
Chain 4:
Chain 4: Iteration:
                     1 / 15000 [ 0%] (Warmup)
Chain 4: Iteration:
                  1500 / 15000 [ 10%] (Warmup)
Chain 4: Iteration:
                  3000 / 15000 [ 20%] (Warmup)
Chain 4: Iteration: 4500 / 15000 [ 30%] (Warmup)
Chain 4: Iteration: 5001 / 15000 [ 33%] (Sampling)
Chain 4: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 4: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 4: Iteration: 9500 / 15000 [ 63%] (Sampling)
Chain 4: Iteration: 11000 / 15000 [ 73%] (Sampling)
Chain 4: Iteration: 12500 / 15000 [ 83%]
                                        (Sampling)
Chain 4: Iteration: 14000 / 15000 [ 93%]
                                        (Sampling)
```

```
Chain 4: Iteration: 15000 / 15000 [100%] (Sampling)
Chain 4.
Chain 4:
        Elapsed Time: 0.06 seconds (Warm-up)
Chain 4:
                       0.124 seconds (Sampling)
Chain 4:
                       0.184 seconds (Total)
Chain 4:
# 사후표본 추출
posterior samples <- extract(fit)</pre>
print(posterior samples)
$theta
  [1] 3.445603 5.207336 4.254783 3.819930 3.201131 3.595814 3.273514 3.407784
3.827176 4.339007 3.603158 2.978973
 [13] 3.440691 4.423264 4.107595 3.783178 4.716802 3.142380 4.762579 4.096370
4.276408 5.372437 3.899219 4.122407
 [25] 4.392978 4.366621 4.104879 4.122928 3.960988 3.222817 3.637218 3.349401
4.281940 4.370535 3.663038 3.960539
 [37] 4.171915 3.649671 3.542375 4.077354 4.516309 5.250884 4.567541 4.032664
3.548034 4.024496 3.767693 3.295317
 [49] 3.814808 3.866771 4.781169 3.921401 3.228510 2.674731 4.314803 2.844673
4.361176 3.744820 4.071879 3.710208
 1611 3.357638 3.818976 5.493548 4.681274 4.239540 3.782264 3.301155 3.587947
5.154232 3.341497 3.896728 4.089677
 [73] 4.216479 4.175259 4.819318 4.358551 3.627441 5.300617 3.146992 3.872979
6.105203 4.731770 3.290179 4.085995
 [85] 3.828614 3.631023 3.540945 3.997197 3.559003 3.574714 3.562595 5.554293
3.455027 3.573720 4.268640 4.524286
 [97] 2.669963 2.810806 4.587997 4.489633 3.845256 3.726410 3.564610 3.171271
4.180055 3.455291 4.487555 3.739379
[109] 2.988388 4.699321 3.608142 4.213024 4.073908 3.531686 3.218229 3.903656
3.748002 3.945296 3.408384 3.525859
[121] 4.314633 2.855916 4.005181 3.954305 3.435033 4.982527 3.105177 3.818553
2.999924 4.151980 4.326920 5.135261
[133] 3.547843 3.667110 3.850102 4.269439 3.696120 4.392461 3.574146 4.054863
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```

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3.762157 3.830322 3.642266 4.758303
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```

```
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3.238146 4.267006 3.426818 5.034179
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[ reached getOption("max.print") -- omitted 39000 entries ]
```

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13.405041 11.226354 11.841548 13.316369		
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13.452624 13.776649 12.476120 13.143297		
[111] 13.685954 13.721448 13.794962 13.609229 13.091397	13.821898	13.780864
13.822009 13.446059 13.602631 13.640379		
[122] 12.017001 13.814476 13.821451 13.485586 12.605580	12.815946	13.807581
12.512308 13.759179 13.629072 12.261754	10 500 10 1	10.65.1077
[133] 13.626961 13.732957 13.815133 13.679202 13.752308	13.563424	13.6540//
13.801512 13.192529 13.670182 13.539910	10 101705	10.666766
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13.694370 13.144158 13.630396 13.762022	12.740072	42 5 45 64 6
[155] 13.595430 13.685502 13.772725 13.734631 13.813811	13./408/2	13.545616
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13.504158 12.393656 13.798533 12.237499	
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13.539447 13.239030 13.705854 13.389880	
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[595] 10.945785 13.641724 13.126904 13.692857 10.406077	13.454112	13.609099
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[705] 13.604219 13.300724 13.805169 13.716559 13.705172	13.078557	13.723772
13.366467 13.602080 12.935487 11.184866		
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13.769338 13.822511 13.822449 13.531263		
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13.401738 13.678500 13.811465 13.418541			
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13.326022 10.117975 13.773658 13.811617			
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13.750095 13.821772 13.802697 13.730327			
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11.120157 13.690908 11.708525 13.714790			
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13.508073 12.290036 13.448198 12.542015			
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13.432319 13.243840 12.147567 13.811985			
[914] 12.889752 13.795744 13.146079 13.361415	13.239438	13.821663	12.316775
12.896982 13.439746 13.645402 11.706041			
[925] 13.607625 13.478438 12.795927 13.820436	13.784694	13.438845	13.789853
13.706123 13.687062 13.142393 13.803297			
[936] 13.119456 13.817521 13.686387 13.749497	13.775942	13.657410	13.710853
13.472769 13.781977 12.461498 13.722550			
[947] 13.801773 13.777649 13.151449 13.754352	13.764406	13.460335	13.028005
13.487563 13.812844 12.585879 11.772130	40.040.77	40 -0 :	40 44====
[958] 12.884327 12.709448 13.814054 9.220104	13.819621	13.724537	13.415565
12.228368 13.495420 13.775410 13.047823	40.6.6.=-	40 500= :-	40 =====
[969] 13.362378 11.829364 13.624523 13.728120	13.640876	12.528745	13.555046
13.752061 13.010562 13.790135 12.516391	42.005054	40.770640	12.000005
[980] 13.278287 13.822511 13.718075 13.554693	13.085951	13.//0640	13.806285

12.144592 13.794375 12.399521 13.802995

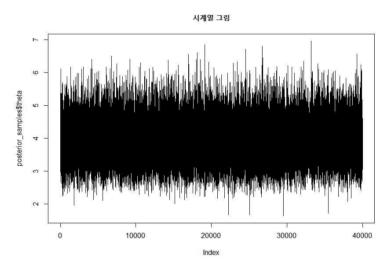
[991] 12.347236 12.689956 13.134759 13.681167 13.473658 12.493335 13.579312 11.892974 13.822099 13.596865

[reached getOption("max.print") -- omitted 39000 entries]

(b)

시계열 그림

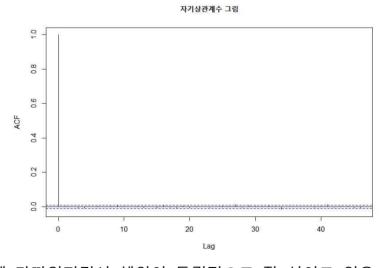
plot(posterior_samples\$theta, type = "l", main = "시계열 그림")



- 각 체인이 3~5 사이의 특정범위에서 집중적으로 변동

자기상관계수 그림

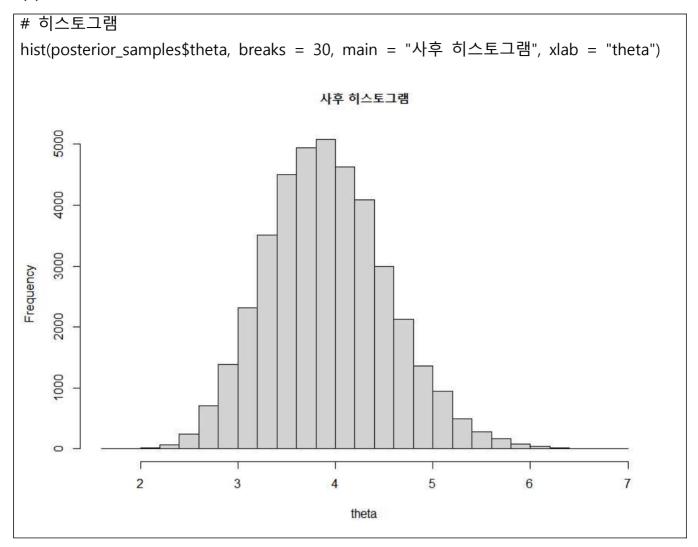
acf(posterior_samples\$theta, main = "자기상관계수 그림")



- 자기상관이 0에 가까워지면서 체인이 독립적으로 잘 섞이고 있음

```
# Stan 결과를 각 체인별로 `mcmc` 객체로 변환
# 체인별 추출 후 'coda::mcmc' 객체로 변환
chains <- extract(fit, permuted = FALSE)</pre>
mcmc chains <- lapply(1:dim(chains)[2], function(chain) mcmc(as.matrix(chains[, chain,
1)))
# 'mcmc.list' 객체로 결합
mcmc list <- mcmc.list(mcmc chains)</pre>
# Gelman-Rubin 진단
gelman_diag <- gelman.diag(mcmc_list)</pre>
print(gelman diag)
Potential scale reduction factors:
     Point est. Upper C.I.
theta
lp__
            1
                      1
Multivariate psrf
1
=> Point estimate와 Upper C.I.가 모두 1: 이는 체인이 수렴했음을 나타냄. 일반적으로
PSRF 값이 1에 가까울수록, 특히 1.1 이하일 때 체인이 수렴했다고 간주할 수 있음.
여기서는 theta와 lp 두 변수 모두 1로 나타나므로 수렴한 것으로 판단할 수 있음.
Multivariate PSRF 값이 1: 다변량 진단에서도 체인들이 수렴했음을 보여줌.
따라서, 이 결과에 따르면 마르코프 체인이 수렴했다고 결론지을 수 있음.
```

(c)



(d)

```
# 통계량 계산
posterior_mean <- mean(posterior_samples$theta)
posterior_sd <- sd(posterior_samples$theta)
credible_interval <- quantile(posterior_samples$theta, probs = c(0.025, 0.975))
print(list(mean = posterior_mean, sd = posterior_sd, Cl = credible_interval))
$mean
[1] 3.920841

$sd
[1] 0.6242959

$CI
2.5% 97.5%
2.789188 5.227749
```

(a)

```
# 필요한 패키지 로드
library(rstan)
library(coda)
# 데이터
data list <- list(
  N = 10,
  x = c(1, 2, 1, 0, 1, 0, 0, 2, 3, 3),
  y = c(0, 1, 0, 0, 0, 2, 1, 2, 3, 1)
# Stan 모델 정의
stan model <- "
data {
  int<lower=0> N;
  int < lower = 0 > x[N];
  int<lower=0> y[N];
}parameters {
  real<lower=0> theta1;
  real<lower=0> theta2;
}model {
  theta1 ~ exponential(1);
  theta2 ~ exponential(1);
  x \sim poisson(theta1);
  y ~ poisson(theta2);
}generated quantities {
  real diff = theta1 - theta2;
# Stan 모델 컴파일 및 샘플링
fit <- stan(model_code = stan_model, data = data_list, iter = 15000, warmup = 5000,
chains = 4)
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 1).
Chain 1:
Chain 1: Gradient evaluation took 2.6e-05 seconds
Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.26
seconds.
Chain 1: Adjust your expectations accordingly!
```

```
Chain 1:
Chain 1:
Chain 1: Iteration:
                      1 / 15000 [ 0%]
                                       (Warmup)
Chain 1: Iteration:
                  1500 / 15000 [ 10%] (Warmup)
Chain 1: Iteration:
                  3000 / 15000 [ 20%] (Warmup)
Chain 1: Iteration:
                  4500 / 15000 [ 30%] (Warmup)
Chain 1: Iteration: 5001 / 15000 [ 33%] (Sampling)
Chain 1: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 1: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 1: Iteration: 9500 / 15000 [ 63%] (Sampling)
Chain 1: Iteration: 11000 / 15000 [ 73%] (Sampling)
Chain 1: Iteration: 12500 / 15000 [ 83%] (Sampling)
Chain 1: Iteration: 14000 / 15000 [ 93%]
                                         (Sampling)
Chain 1: Iteration: 15000 / 15000 [100%]
                                         (Sampling)
Chain 1:
Chain 1:
         Elapsed Time: 0.072 seconds (Warm-up)
Chain 1:
                        0.161 seconds (Sampling)
Chain 1.
                        0.233 seconds (Total)
Chain 1:
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 2).
Chain 2:
Chain 2: Gradient evaluation took 4e-06 seconds
Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.04
seconds.
Chain 2: Adjust your expectations accordingly!
Chain 2:
Chain 2:
Chain 2: Iteration:
                      1 / 15000 [ 0%]
                                       (Warmup)
Chain 2: Iteration:
                  1500 / 15000 [ 10%] (Warmup)
Chain 2: Iteration:
                  3000 / 15000 [ 20%] (Warmup)
Chain 2: Iteration: 4500 / 15000 [ 30%] (Warmup)
Chain 2: Iteration:
                  5001 / 15000 [ 33%] (Sampling)
Chain 2: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 2: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 2: Iteration: 9500 / 15000 [ 63%]
                                        (Sampling)
Chain 2: Iteration: 11000 / 15000 [ 73%]
                                         (Sampling)
Chain 2: Iteration: 12500 / 15000 [ 83%]
                                         (Sampling)
```

```
Chain 2: Iteration: 14000 / 15000 [ 93%] (Sampling)
Chain 2: Iteration: 15000 / 15000 [100%] (Sampling)
Chain 2:
Chain 2: Elapsed Time: 0.083 seconds (Warm-up)
Chain 2:
                       0.22 seconds (Sampling)
Chain 2:
                       0.303 seconds (Total)
Chain 2:
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 3).
Chain 3:
Chain 3: Gradient evaluation took 6e-06 seconds
Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.06
seconds.
Chain 3: Adjust your expectations accordingly!
Chain 3:
Chain 3:
Chain 3: Iteration:
                     1 / 15000 [ 0%] (Warmup)
Chain 3: Iteration: 1500 / 15000 [ 10%] (Warmup)
Chain 3: Iteration: 3000 / 15000 [ 20%] (Warmup)
Chain 3: Iteration: 4500 / 15000 [ 30%] (Warmup)
Chain 3: Iteration: 5001 / 15000 [ 33%] (Sampling)
Chain 3: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 3: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 3: Iteration: 9500 / 15000 [ 63%] (Sampling)
Chain 3: Iteration: 11000 / 15000 [ 73%] (Sampling)
Chain 3: Iteration: 12500 / 15000 [ 83%] (Sampling)
Chain 3: Iteration: 14000 / 15000 [ 93%] (Sampling)
Chain 3: Iteration: 15000 / 15000 [100%] (Sampling)
Chain 3:
Chain 3: Elapsed Time: 0.089 seconds (Warm-up)
Chain 3:
                       0.228 seconds (Sampling)
Chain 3:
                       0.317 seconds (Total)
Chain 3:
SAMPLING FOR MODEL 'anon model' NOW (CHAIN 4).
Chain 4:
Chain 4: Gradient evaluation took 5e-06 seconds
Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.05
```

```
seconds.
Chain 4: Adjust your expectations accordingly!
Chain 4:
Chain 4:
Chain 4: Iteration:
                     1 / 15000 [ 0%] (Warmup)
Chain 4: Iteration:
                 1500 / 15000 [ 10%] (Warmup)
Chain 4: Iteration:
                 3000 / 15000 [ 20%] (Warmup)
Chain 4: Iteration: 4500 / 15000 [ 30%] (Warmup)
Chain 4: Iteration: 5001 / 15000 [ 33%] (Sampling)
Chain 4: Iteration: 6500 / 15000 [ 43%] (Sampling)
Chain 4: Iteration: 8000 / 15000 [ 53%] (Sampling)
Chain 4: Iteration: 9500 / 15000 [ 63%] (Sampling)
Chain 4: Iteration: 11000 / 15000 [ 73%] (Sampling)
Chain 4: Iteration: 12500 / 15000 [ 83%] (Sampling)
Chain 4: Iteration: 14000 / 15000 [ 93%] (Sampling)
Chain 4: Iteration: 15000 / 15000 [100%]
                                        (Sampling)
Chain 4:
Chain 4:
         Elapsed Time: 0.093 seconds (Warm-up)
Chain 4:
                       0.182 seconds (Sampling)
Chain 4:
                       0.275 seconds (Total)
Chain 4:
# 사후표본 추출
posterior samples <- extract(fit)
print(posterior_samples)
$theta1
  [1] 1.2668631 0.9731773 1.8738613 1.6145547 1.8354870 0.8144773 1.5930936
0.7378882 1.3575845 1.3410056 1.1689440
 [12] 1.2336804 1.1203088 1.6578888 1.0450051 1.1271575 0.9089330 1.1546596
1.2478028 1.3841022 1.2977756 2.0417244
 [23] 1.0152349 1.1265221 1.0880690 1.0277726 1.5290695 1.0362229
                                                                        1.2778133
0.7360973 1.6959353 1.6413268 1.0552767
 [34] 1.0407748 1.7079826 0.8244508 1.1569542 1.3878469 1.2206447
                                                                        1.2374660
1.7206365 1.2977368 1.0812016 1.5701236
 [45] 0.7633375 1.8481562 0.9337708 1.4867301 1.5938793 0.8216459
                                                                        1.0162456
1.3131414 1.7944806 1.6038961 1.2434235
  [56] 1.2205768 1.0368313 0.9738308 1.7835393 1.0649060 0.9742341 1.2951844
```

1.9752971 2.0535916 1.4725410 0.8367019 [67] 0.9020565 1.3810321 0.6918161 1.1021510 1.3699514 1.2783553 1.3172772 0.8546070 0.7692252 0.9822657 1.7344512 0.9853500 1.2044925 0.8584834 1.1673813 0.9761475 1.5254130 1.6402381 0.8139369 0.7643350 1.7047254 1.4658326 1.0674242 1.3756989 0.9269461 0.8047735 1.1094800 0.8640304 1.2988338 1.00] 1.6831902 1.4718982 1.3873373 0.9438492 1.5471149 0.6305330 1.1750871 0.8556096 1.0016541 0.7286500 1.6538655 1.11] 1.1725288 1.3147777 1.7603476 1.3565938 1.5524007 0.8971626 1.1439511 1.4989646 1.5879278 1.8256709 0.8920750 1.22] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.432
0.8546070 0.7692252 0.9822657 1.7344512 [78] 0.9756285 0.7837877 1.2926695 0.9921817 0.9853500 1.2044925 0.8584834 1.1673813 0.9761475 1.5254130 1.6402381 [89] 0.8139369 0.7643350 1.7047254 1.4658326 1.0674242 1.3756989 0.9269461 0.8047735 1.1094800 0.8640304 1.2988338 [100] 1.6831902 1.4718982 1.3873373 0.9438492 1.5471149 0.6305330 1.1750871 0.8556096 1.0016541 0.7286500 1.6538655 [111] 1.1725288 1.3147777 1.7603476 1.3565938 1.5524007 0.8971626 1.1439511 1.4989646 1.5879278 1.8256709 0.8920750 [122] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 [133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.4322828 1.0182181 0.6677236
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1.1673813 0.9761475 1.5254130 1.6402381 [89] 0.8139369 0.7643350 1.7047254 1.4658326 1.0674242 1.3756989 0.9269461 0.8047735 1.1094800 0.8640304 1.2988338 1.010] 1.6831902 1.4718982 1.3873373 0.9438492 1.5471149 0.6305330 1.1750871 0.8556096 1.0016541 0.7286500 1.6538655 [111] 1.1725288 1.3147777 1.7603476 1.3565938 1.5524007 0.8971626 1.1439511 1.4989646 1.5879278 1.8256709 0.8920750 [122] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 [133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.4322828 1.0182181 0.6677236
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0.8047735 1.1094800 0.8640304 1.2988338 [100] 1.6831902 1.4718982 1.3873373 0.9438492 1.5471149 0.6305330 1.1750871 0.8556096 1.0016541 0.7286500 1.6538655 [111] 1.1725288 1.3147777 1.7603476 1.3565938 1.5524007 0.8971626 1.1439511 1.4989646 1.5879278 1.8256709 0.8920750 [122] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 [133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.4322828 1.0182181 0.6677236
[100] 1.6831902 1.4718982 1.3873373 0.9438492 1.5471149 0.6305330 1.1750871 0.8556096 1.0016541 0.7286500 1.6538655 [111] 1.1725288 1.3147777 1.7603476 1.3565938 1.5524007 0.8971626 1.1439511 1.4989646 1.5879278 1.8256709 0.8920750 [122] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 [133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.4322828 1.0182181 0.6677236
0.8556096 1.0016541 0.7286500 1.6538655 [111] 1.1725288 1.3147777 1.7603476 1.3565938 1.5524007 0.8971626 1.1439511 1.4989646 1.5879278 1.8256709 0.8920750 [122] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 [133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.4322828 1.0182181 0.6677236
[111] 1.1725288 1.3147777 1.7603476 1.3565938 1.5524007 0.8971626 1.1439511 1.4989646 1.5879278 1.8256709 0.8920750 [122] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 [133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.4322828 1.0182181 0.6677236
1.4989646 1.5879278 1.8256709 0.8920750 [122] 1.2879956 1.0002390 0.8753255 1.3989410 1.6412030 1.8073886 1.1609687 1.0853394 1.3899120 1.1355043 1.0317581 [133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.5333905 0.8934996 1.4322828 1.0182181 0.6677236
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[133] 0.8464954 1.1497645 1.5102181 1.1541059 1.1583752 1.3449541 1.53333905 0.8934996 1.4322828 1.0182181 0.6677236
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1.8131072 0.9476348 1.0763501 1.2962483		
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1.6328843 1.2995451 1.1821345 1.0512697		
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0.9661888 1.2296188 1.5008860 1.7394893		
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0.7684411 1.9536925 1.4943866 1.2471226		
[617] 1.2863924 1.2059110 1.2260086 1.0104520 1.6681590	0.7292368	1.4768006
1.3947618 1.1319598 0.8570935 1.4132976		
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[585] 0.977821345 0.588936833 0.550218135 0.495737506 -0.399624127 -0.178188110 -0.135327909 -0.024739084 [593] 0.807910403 -0.108645643 0.459648185 0.348748266 0.912448322 0.701305881 -0.299887379 0.297921061 [601] 0.533856019 -0.005356541 0.505476006 0.087103804 1.222861566 0.646179313 -0.339486700 -0.468780056 [609] 0.036216265 0.419869789 0.230315548 1.067516635 -0.901466570 0.943842079 0.147106576 0.466819202 [617] -0.149972174 -0.523815290 -0.574888969 -0.065710090 0.508513009 -0.423741951 0.307275732 0.583043464 [625] 0.191504006 -0.607800773 0.461431778 0.316999581 0.209560634 0.315069159 -0.018387653 0.581580408 [633] 0.010262414 0.890529205 0.436321735 0.190033723 0.551240976 1.290218153 0.018943905 0.889167471 [641] 0.597124541 0.261680067 0.074515487 0.189753327 0.541564654 -0.246632367 0.025902702 -0.363104137 [649] 0.129517294 0.457776380 0.870172879 -0.240655868 -0.231547701 0.742658330 -0.146104641 0.331659245 [657] -0.437210823 0.575872674 0.234705120 0.039435487 0.481873236 0.217925882 0.271363253 0.156256199 [665] 0.279066156 0.275233038 -0.013159416 -0.229976890 0.068275400 -0.755211480 0.474391924 0.062541153 [673] -0.157509720 0.098666451 0.485597076 -0.052373880 0.572658311 0.984732880 -0.144914518 0.252346643	[577] 0.503857540 0.282511986 0.611101492 0.797096169	0.448257256
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[617] -0.149972174 -0.523815290 -0.574888969 -0.065710090 0.508513009 -0.423741951 0.307275732 0.583043464 [625] 0.191504006 -0.607800773 0.461431778 0.316999581 0.209560634 0.315069159 -0.018387653 0.581580408 [633] 0.010262414 0.890529205 0.436321735 0.190033723 0.551240976 1.290218153 0.018943905 0.889167471 [641] 0.597124541 0.261680067 0.074515487 0.189753327 0.541564654 -0.246632367 0.025902702 -0.363104137 [649] 0.129517294 0.457776380 0.870172879 -0.240655868 -0.231547701 0.742658330 -0.146104641 0.331659245 [657] -0.437210823 0.575872674 0.234705120 0.039435487 0.481873236 0.217925882 0.271363253 0.156256199 [665] 0.279066156 0.275233038 -0.013159416 -0.229976890 0.068275400 -0.755211480 0.474391924 0.062541153 [673] -0.157509720 0.098666451 0.485597076 -0.052373880 0.572658311 0.984732880 -0.144914518 0.2523446643	[609] 0.036216265 0.419869789 0.230315548 1.067516635	5 -0.901466570
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[689] 0.817739922 0.514526485 0.544731647 0.159777827 0.129977916	[689] 0.817739922 0.514526485 0.544731647 0.159777827	0.129977916

1.009103905 - 0.077965890 - 0.399270745 [697]	1.000103000 0.077060000 0.300370745		1
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0.192845100		0.080743868	0 996878222
[833] 1.172980866 -0.403563954 -0.236452071 0.488199664 0.829539749 0.220537308 -0.469230309 0.499248206		0.0007 10000	0.550070222
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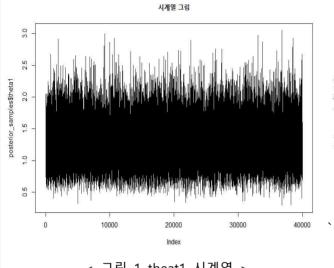
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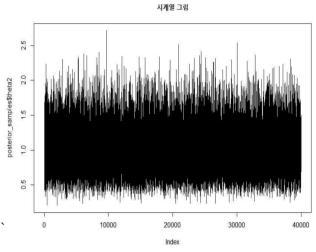
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시계열 그림

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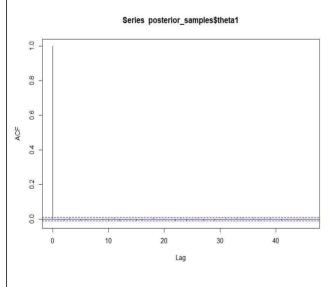


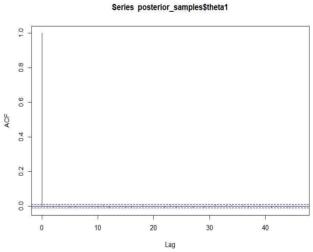
- < 그림 1 theat1 시계열 > < 그림 1 theat1 시계열 >
- (그림 1) 각 체인이 0.5~2 사이의 특정범위에서 집중적으로 변동
- (그림 2) 각 체인이 0.5~1.7 사이의 특정범위에서 집중적으로 변동

자기상관계수 그림

acf(posterior_samples\$theta1)

acf(posterior_samples\$theta2)

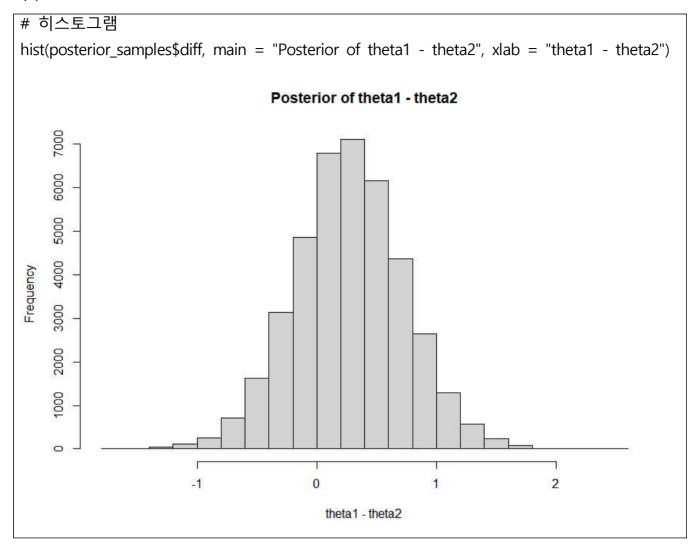




- 자기상관이 0에 가까워지면서 체인이 독립적으로 잘 섞이고 있음

```
# 각 체인을 개별 mcmc 객체로 변환
chains <- extract(fit, permuted = FALSE) # 각 체인을 개별로 추출
# 각 체인을 mcmc 객체로 변환
mcmc chains theta1 <- lapply(1:dim(chains)[2], function(chain) {
 mcmc(as.matrix(chains[, chain, "theta1"]))
})
mcmc_chains_theta2 <- lapply(1:dim(chains)[2], function(chain) {</pre>
  mcmc(as.matrix(chains[, chain, "theta2"]))
})
# mcmc.list로 결합
mcmc list theta1 <- mcmc.list(mcmc chains theta1)
mcmc list theta2 <- mcmc.list(mcmc chains theta2)
# Gelman-Rubin 진단
gelman diag theta1 <- gelman.diag(mcmc list theta1)</pre>
gelman diag theta2 <- gelman.diag(mcmc list theta2)</pre>
print(gelman_diag_theta1)
Potential scale reduction factors:
    Point est. Upper C.I.
[1,]
            1
print(gelman_diag_theta2)
Potential scale reduction factors:
    Point est. Upper C.I.
[1,]
            1
=> Point estimate와 Upper C.I.가 모두 1: 이는 체인이 수렴했음을 나타냄.
여기서는 theta1과 theta2 두 변수 모두 1로 나타나므로 수렴한 것으로 판단할 수 있음.
따라서, 이 결과에 따르면 마르코프 체인이 수렴했다고 결론지을 수 있음.
```

(c)



(d)

```
# 사후평균, 표준편차, 신용구간 계산
mean_diff <- mean(posterior_samples$diff)
sd_diff <- sd(posterior_samples$diff)
ci_diff <- quantile(posterior_samples$diff, probs = c(0.025, 0.975))

# 결과 출력
cat("Posterior mean of theta1 - theta2:", mean_diff, "\n")
Posterior mean of theta1 - theta2: 0.2730707
cat("Posterior standard deviation of theta1 - theta2:", sd_diff, "\n")
Posterior standard deviation of theta1 - theta2: 0.4562278
cat("95% credible interval of theta1 - theta2: -0.6186622 1.179176
```