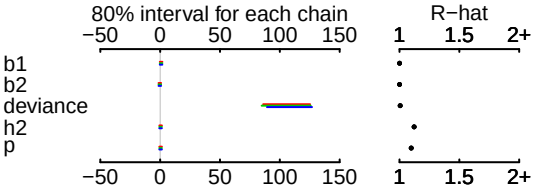
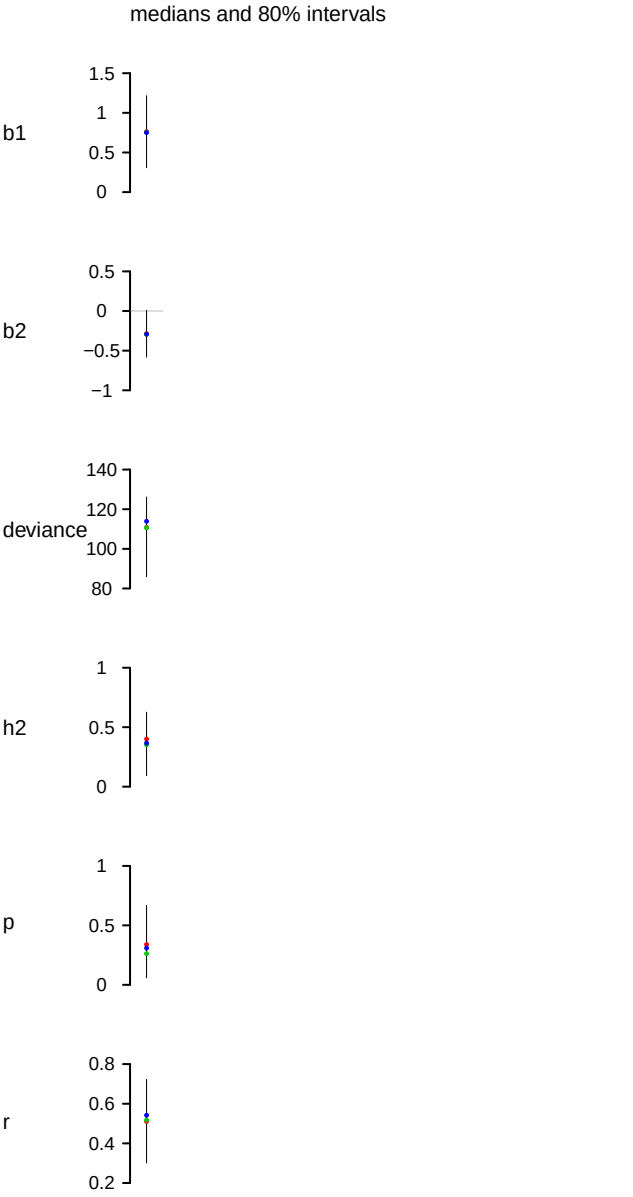
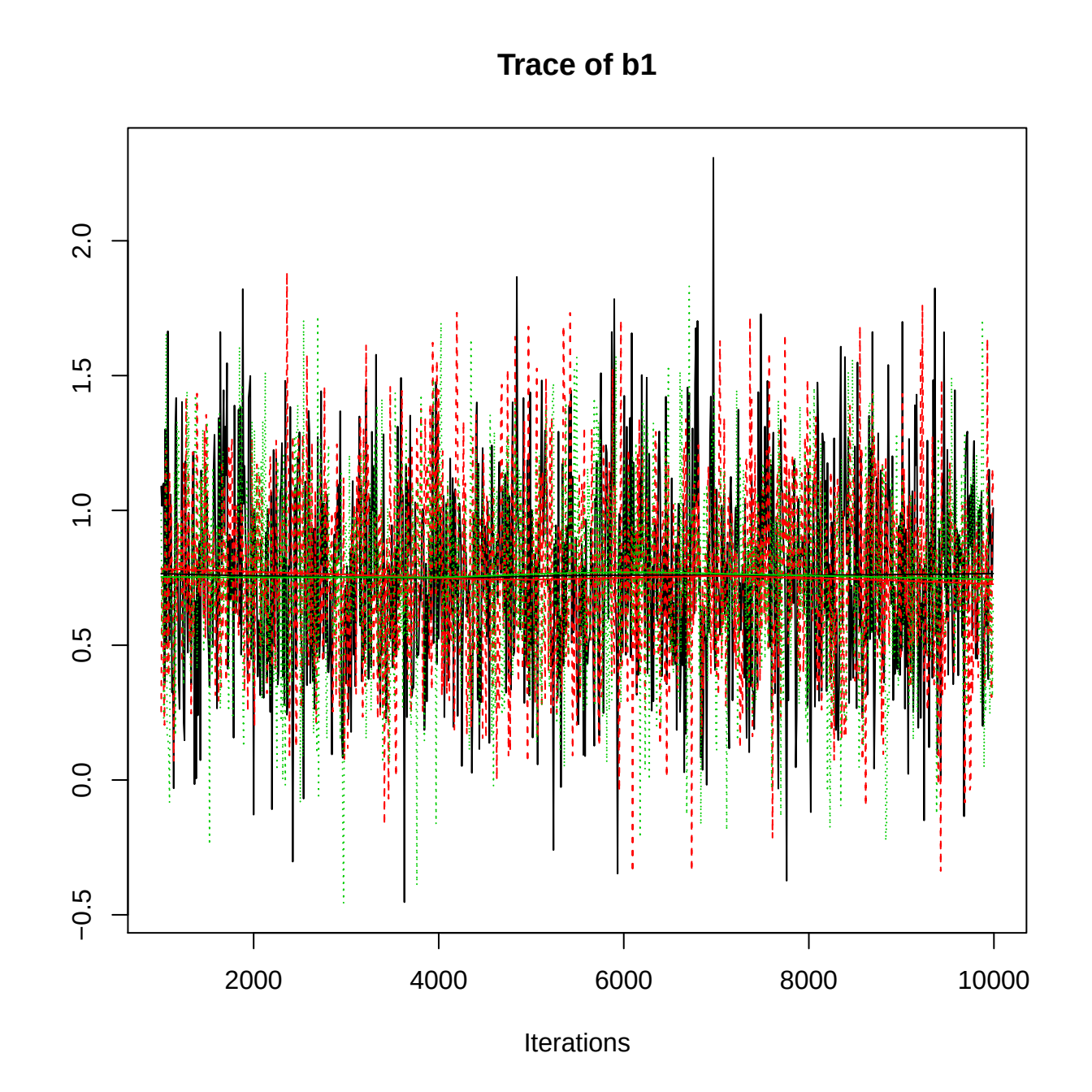
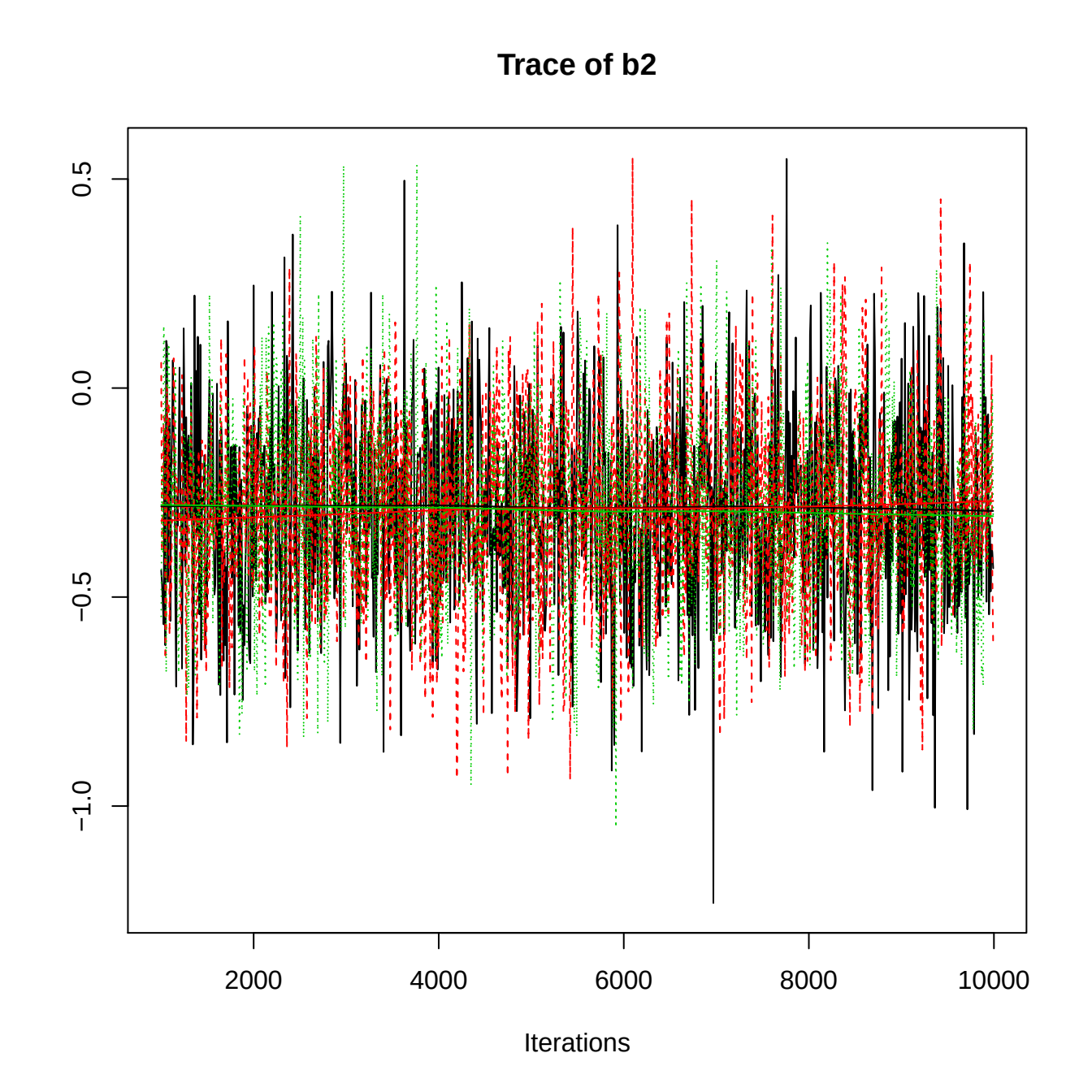
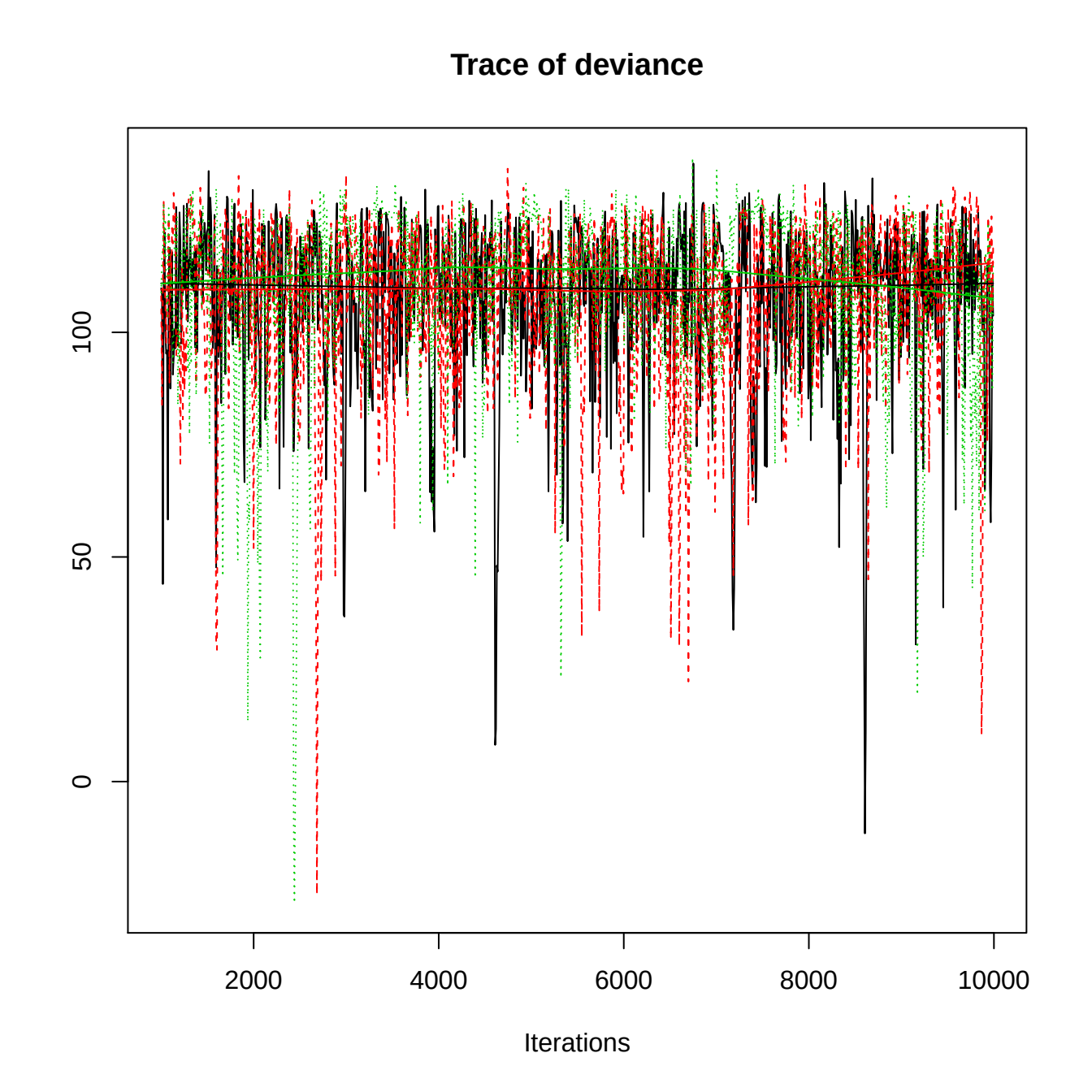
Bugs model at "/tmp/RtmpiIJiSc/model76e225cca161.txt", fit using jags, 3 chains, each with 10000 iterations (first 1000 discarded)

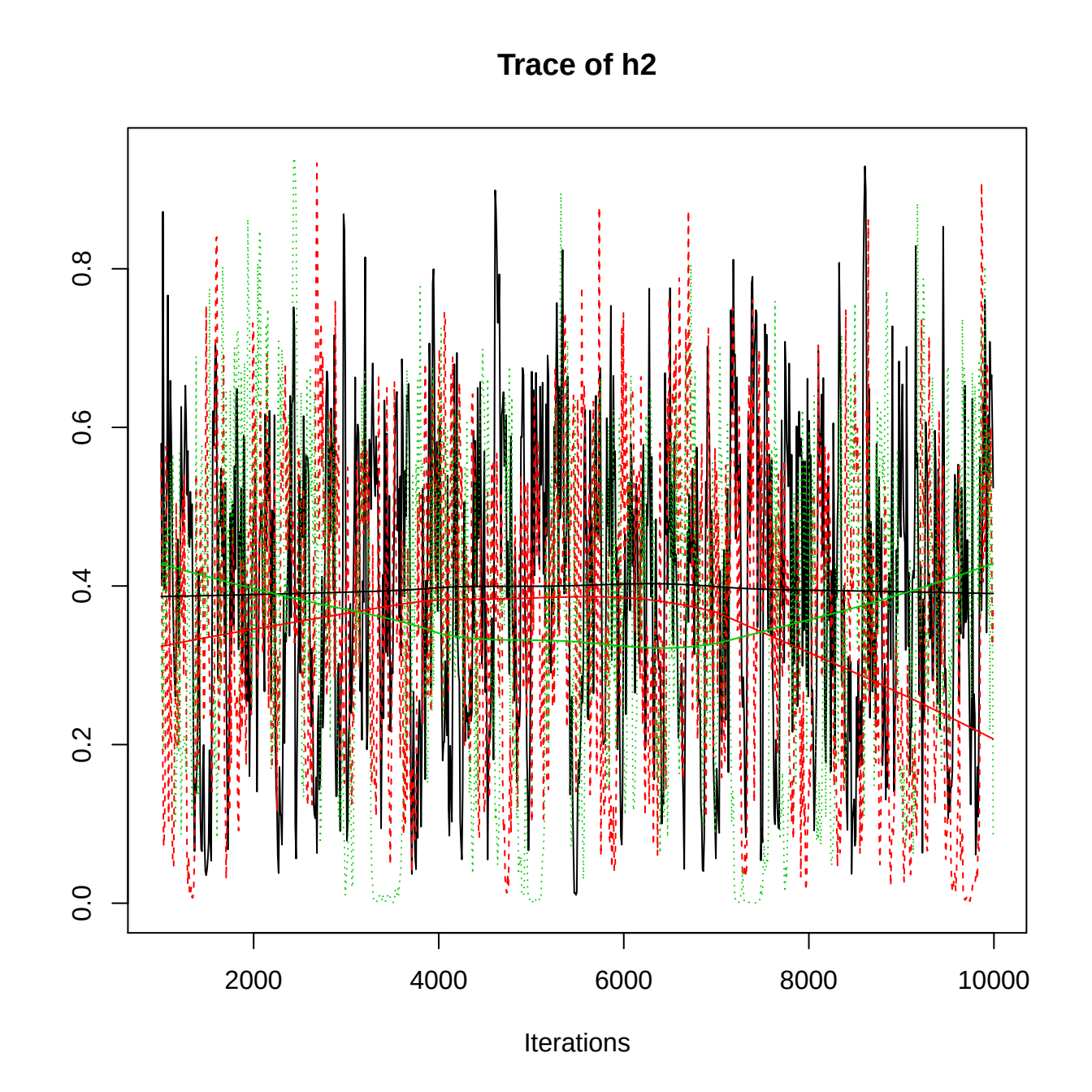


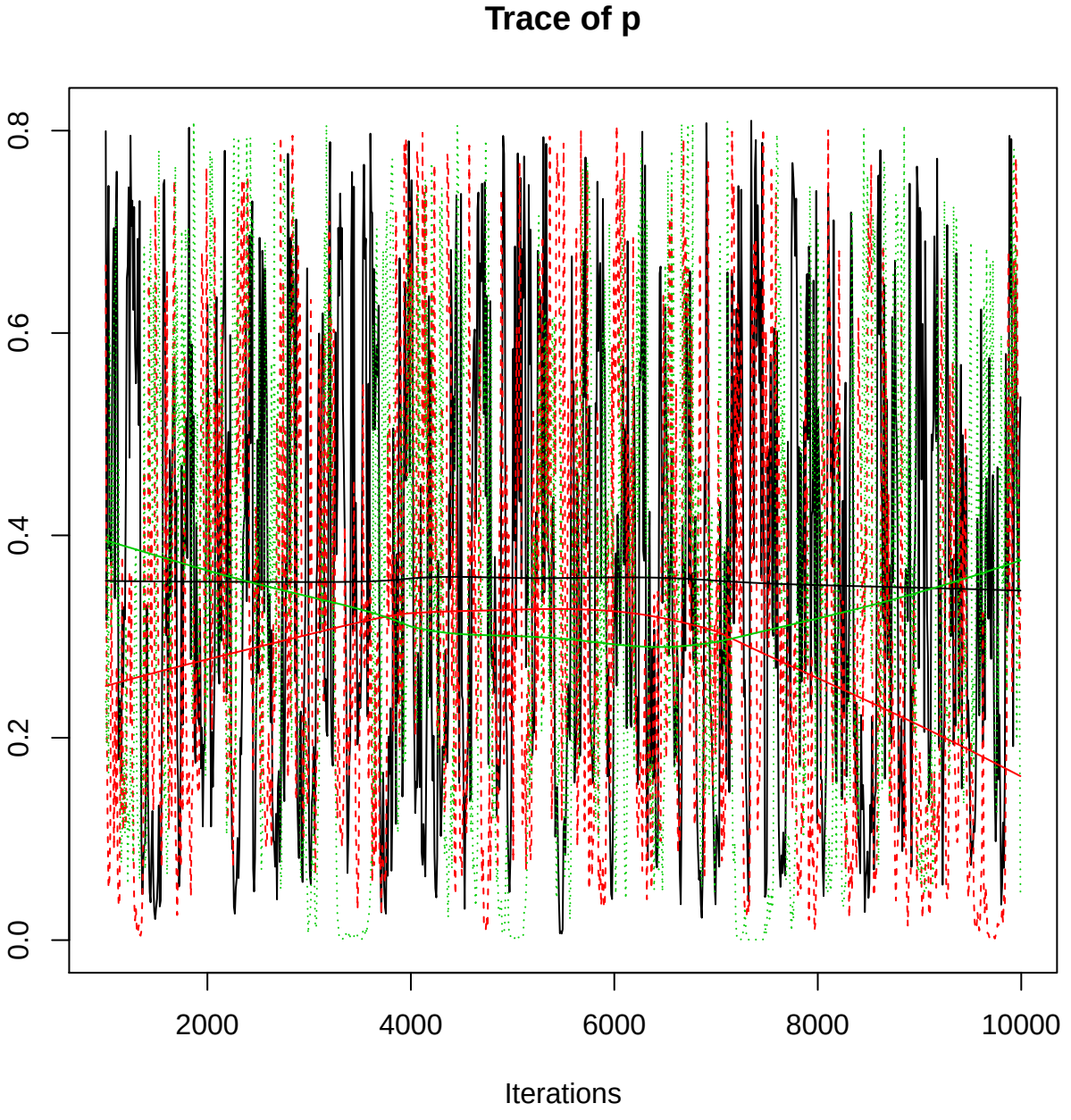


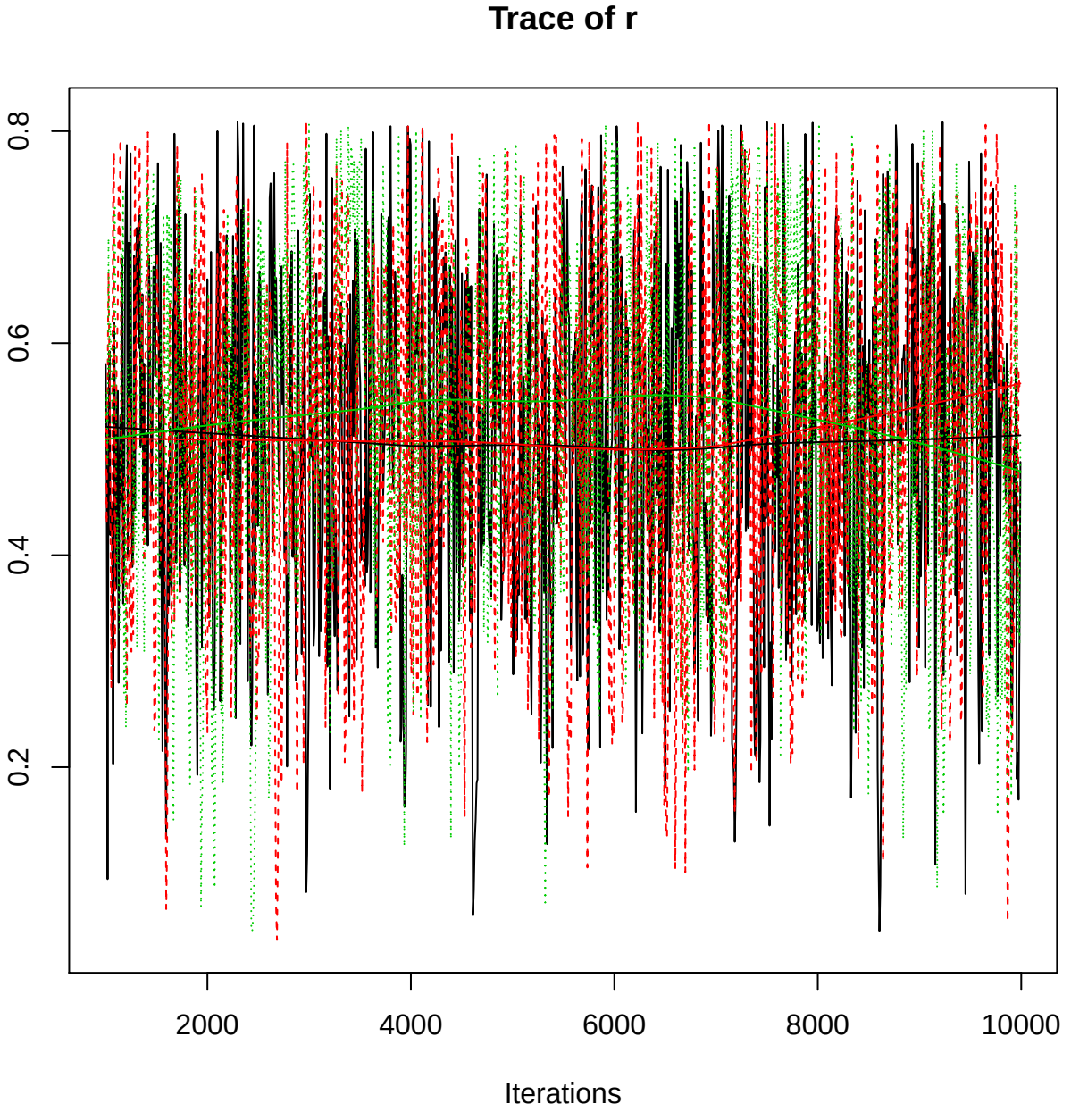

Trace of b1
2.0
1.5
1.0
0.5
0.0
−0.5
2000
4000
6000
8000
10000
Iterations

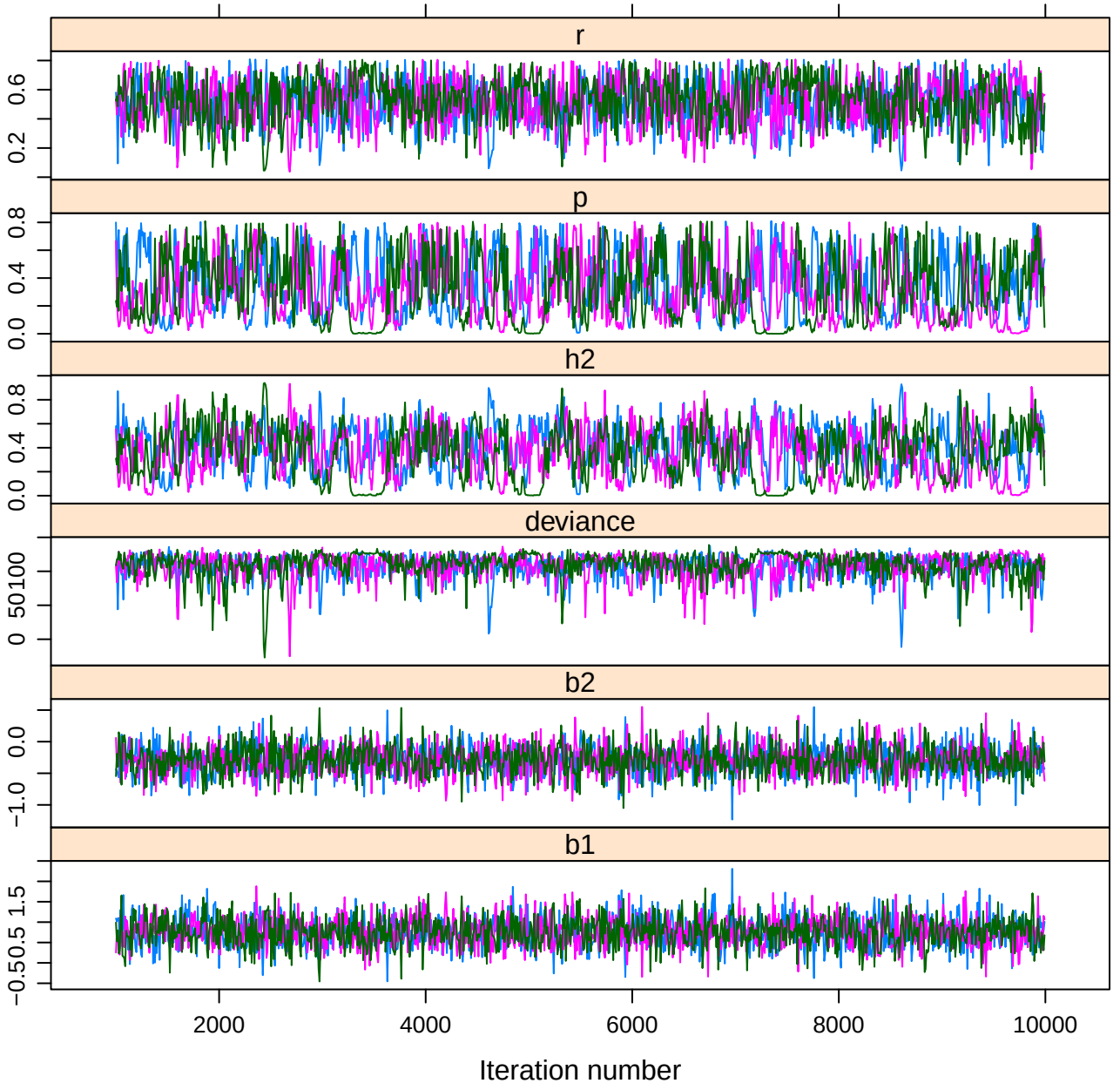

Trace of b2
0.5
0.0
−0.5
−1.0
2000
4000
6000
8000
10000
Iterations

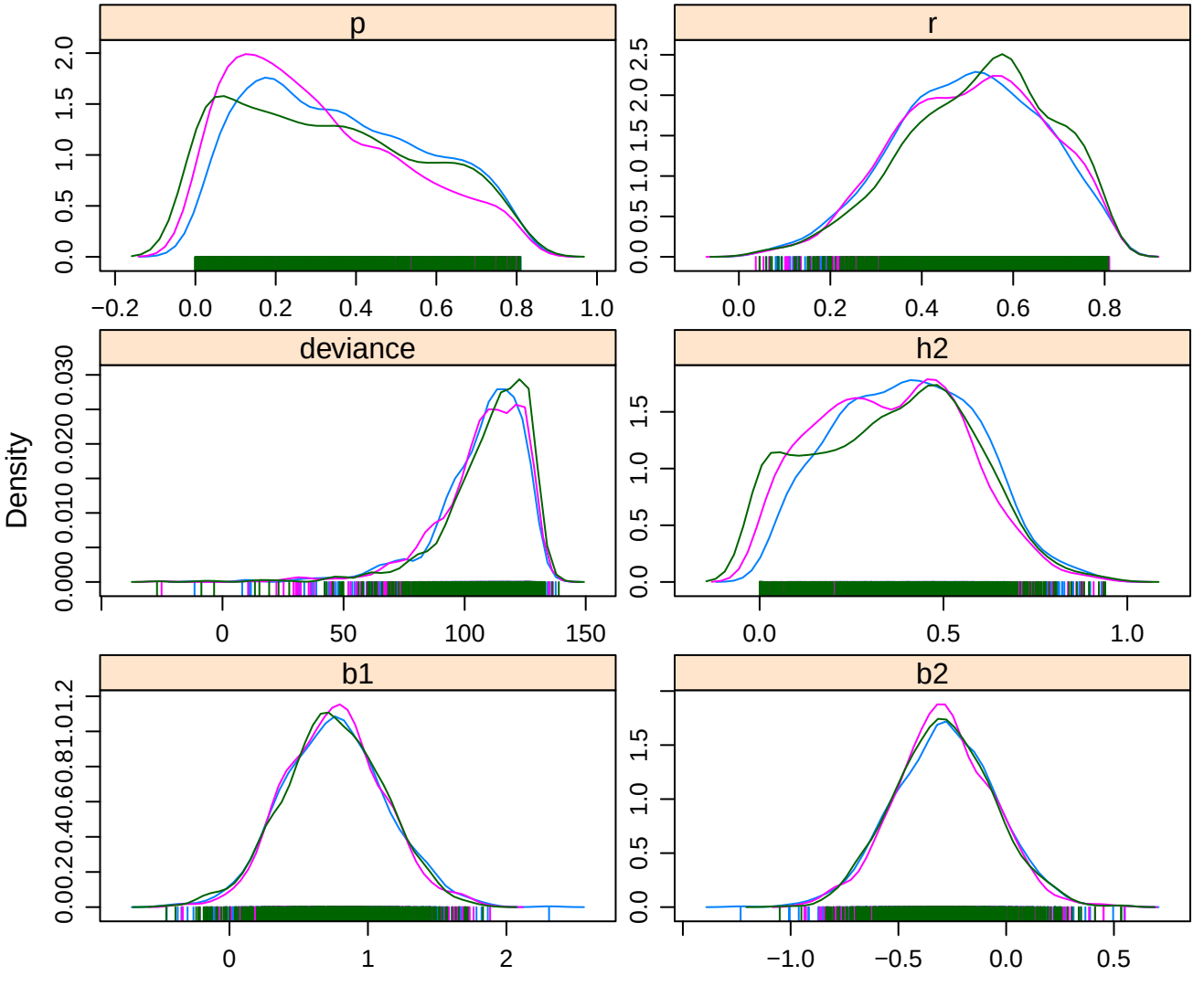

Trace of deviance
0
50
100
2000
4000
6000
8000
10000
Iterations


Trace of h2
0.0
0.2
0.4
0.6
0.8
2000
4000
6000
8000
10000
Iterations


Trace of p
0.0
0.2
0.4
0.6
0.8
2000
4000
6000
8000
10000
Iterations


Trace of r
0.2
0.4
0.6
0.8
2000
4000
6000
8000
10000
Iterations


r
p
h2
deviance
b2
b1
0.2
0.6
0.0
0.4
0.8
0.0
0.4
0.8
0
50
100
−1.0
0.0
−0.5
0.5
1.5
2000
4000
6000
8000
10000
Iteration number


p
r
deviance
h2
b1
b2
Density