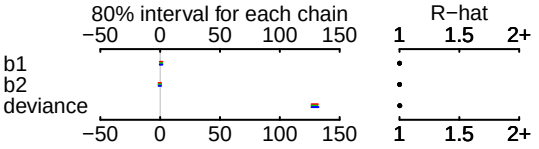
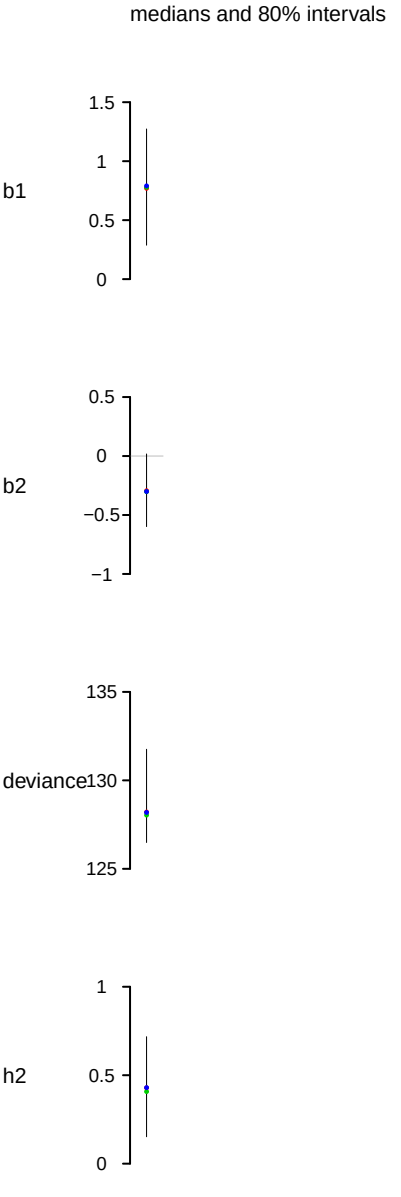
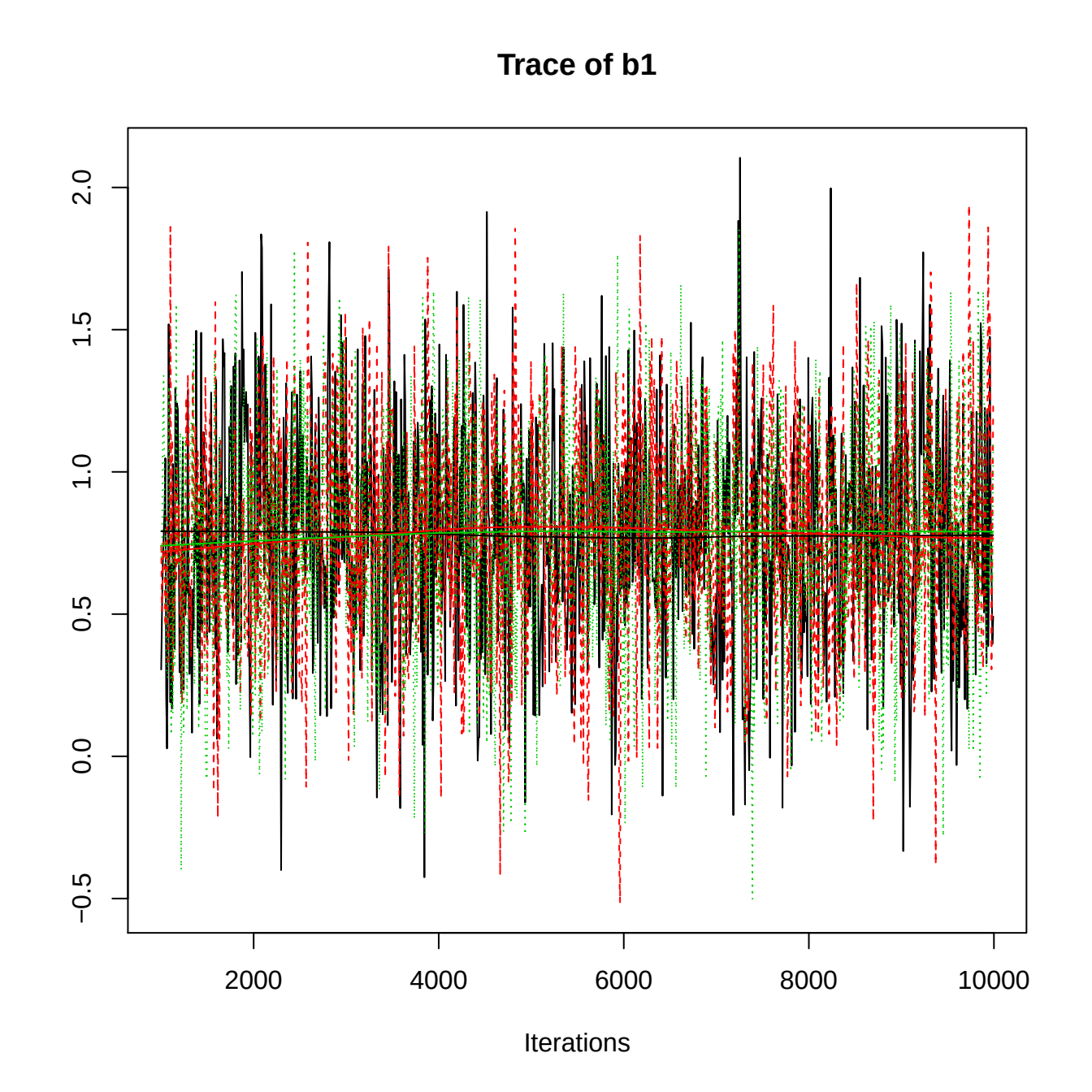
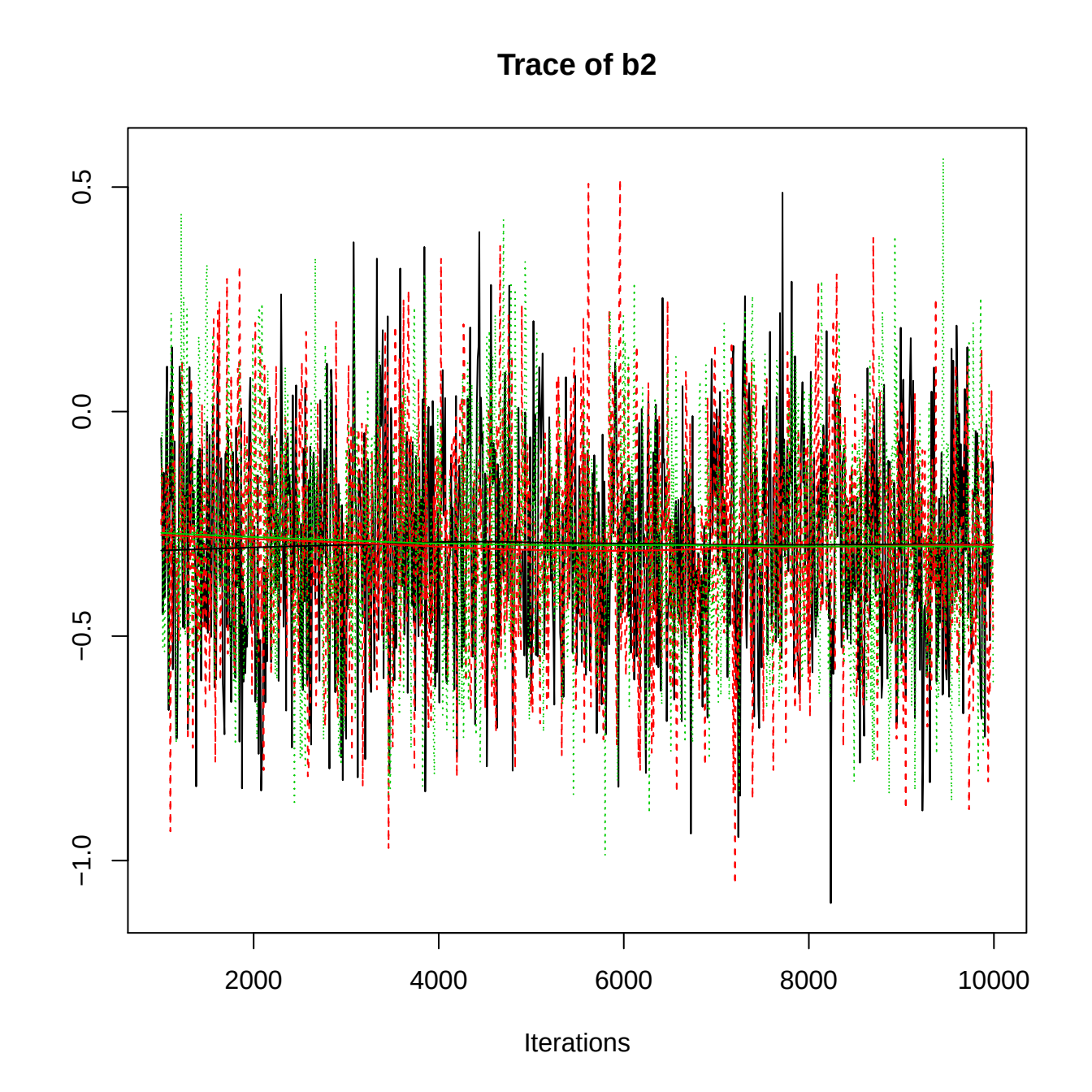
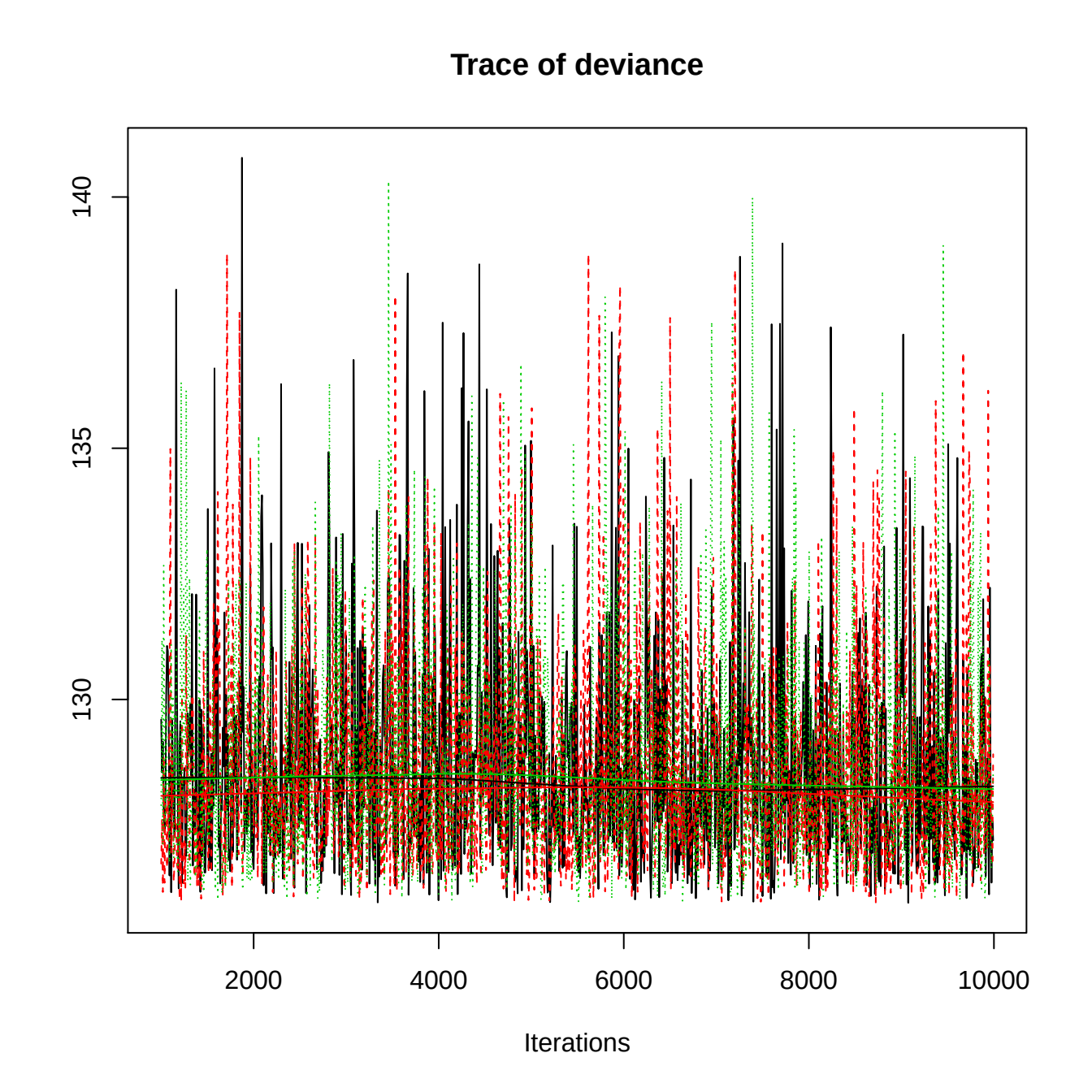
Bugs model at "/tmp/RtmpiIJiSc/model76e22ef3d7c1.txt", fit using jags, 3 chains, each with 10000 iterations (first 1000 discarded)

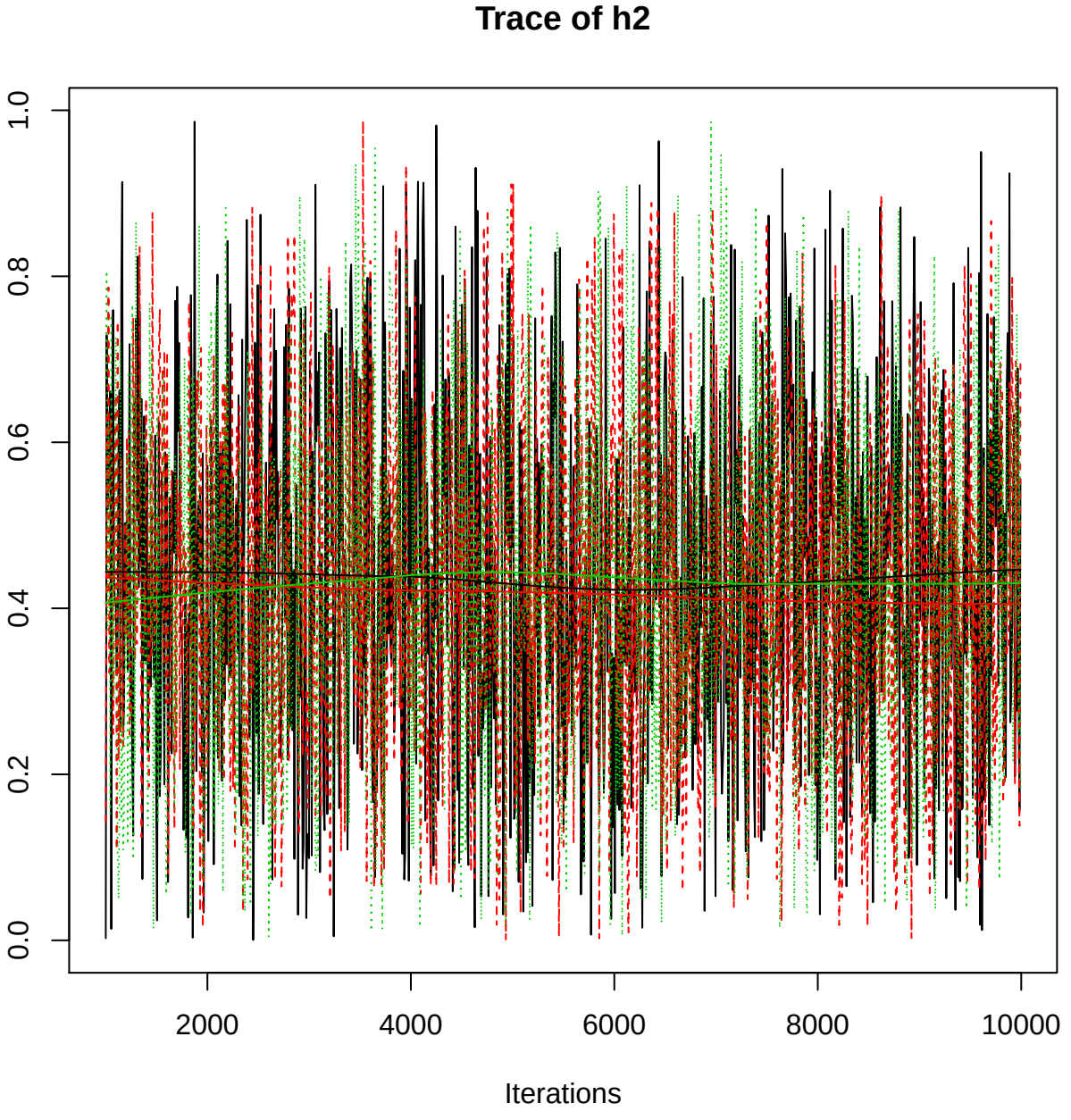


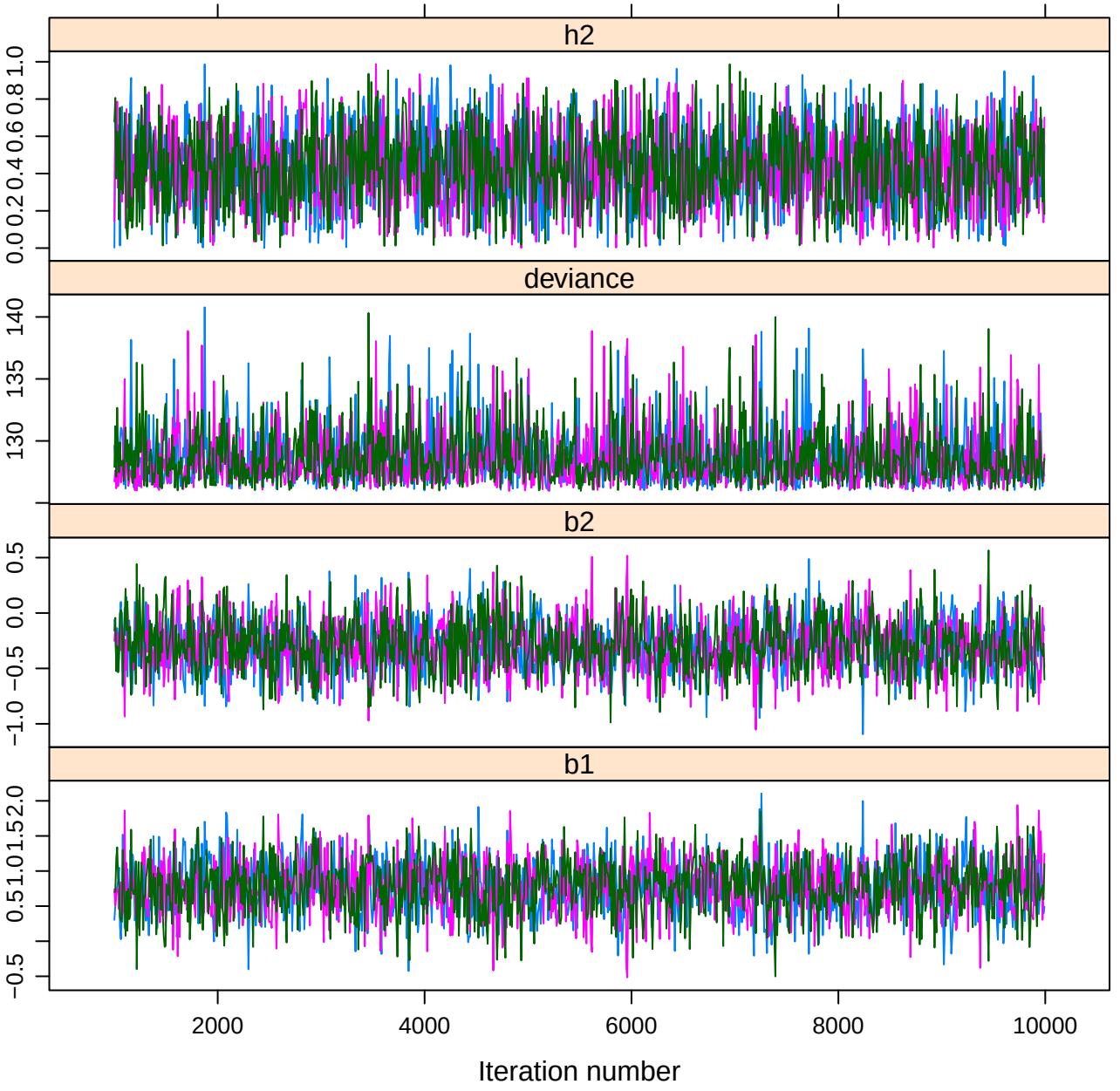


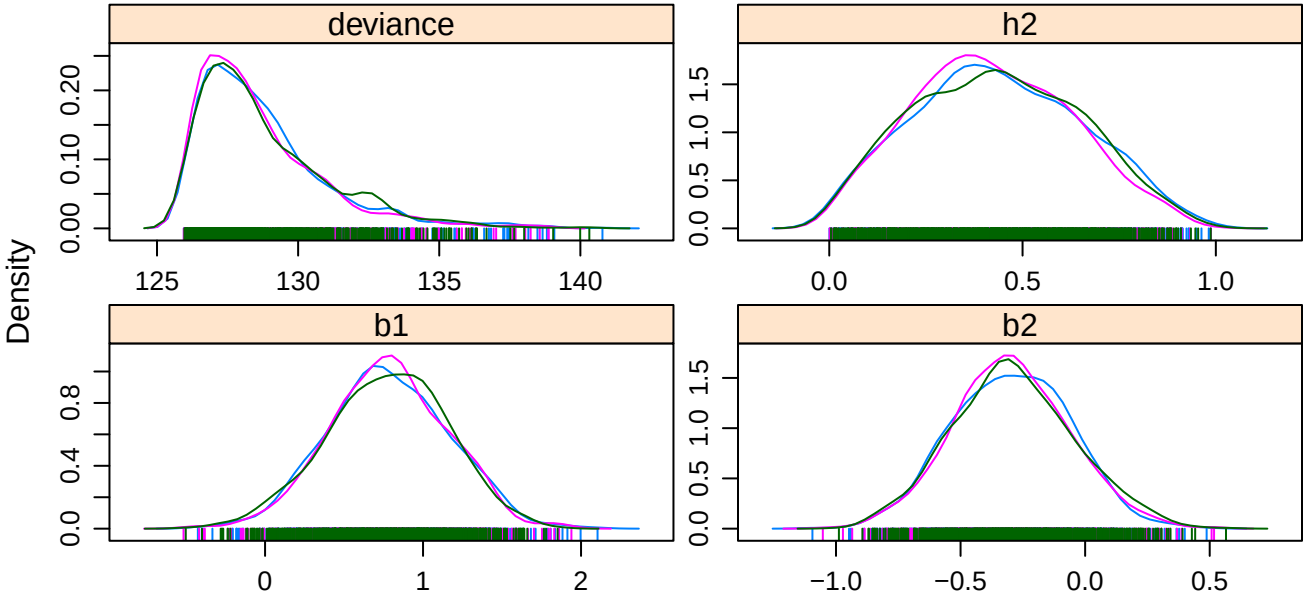

Trace of b1
2.0
1.5
1.0
0.5
0.0
−0.5
2000
4000
6000
8000
10000
Iterations


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


Trace of deviance
140
135
130
2000
4000
6000
8000
10000
Iterations


Trace of h2
1.0
0.8
0.6
0.4
0.2
0.0
2000
4000
6000
8000
10000
Iterations


h2
deviance
b2
b1
Iteration number
2000
4000
6000
8000
10000


deviance
h2
b1
b2
Density
0.00
0.10
0.20
125
130
135
140
0.0
0.5
1.0
1.5
0.0
0.5
1.0
0.0
0.4
0.8
0
1
2
0.0
0.5
1.0
1.5
−1.0
−0.5
0.0
0.5