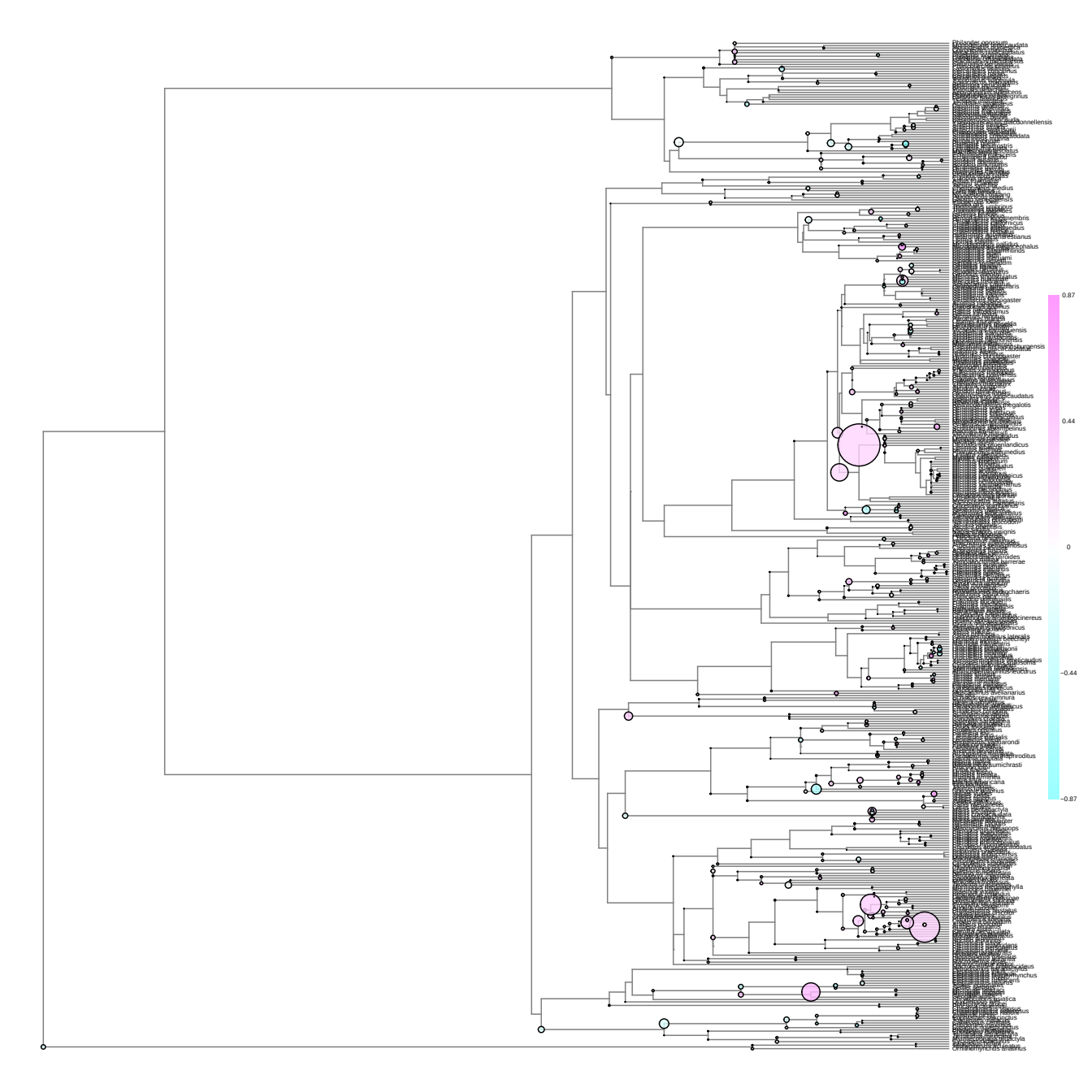
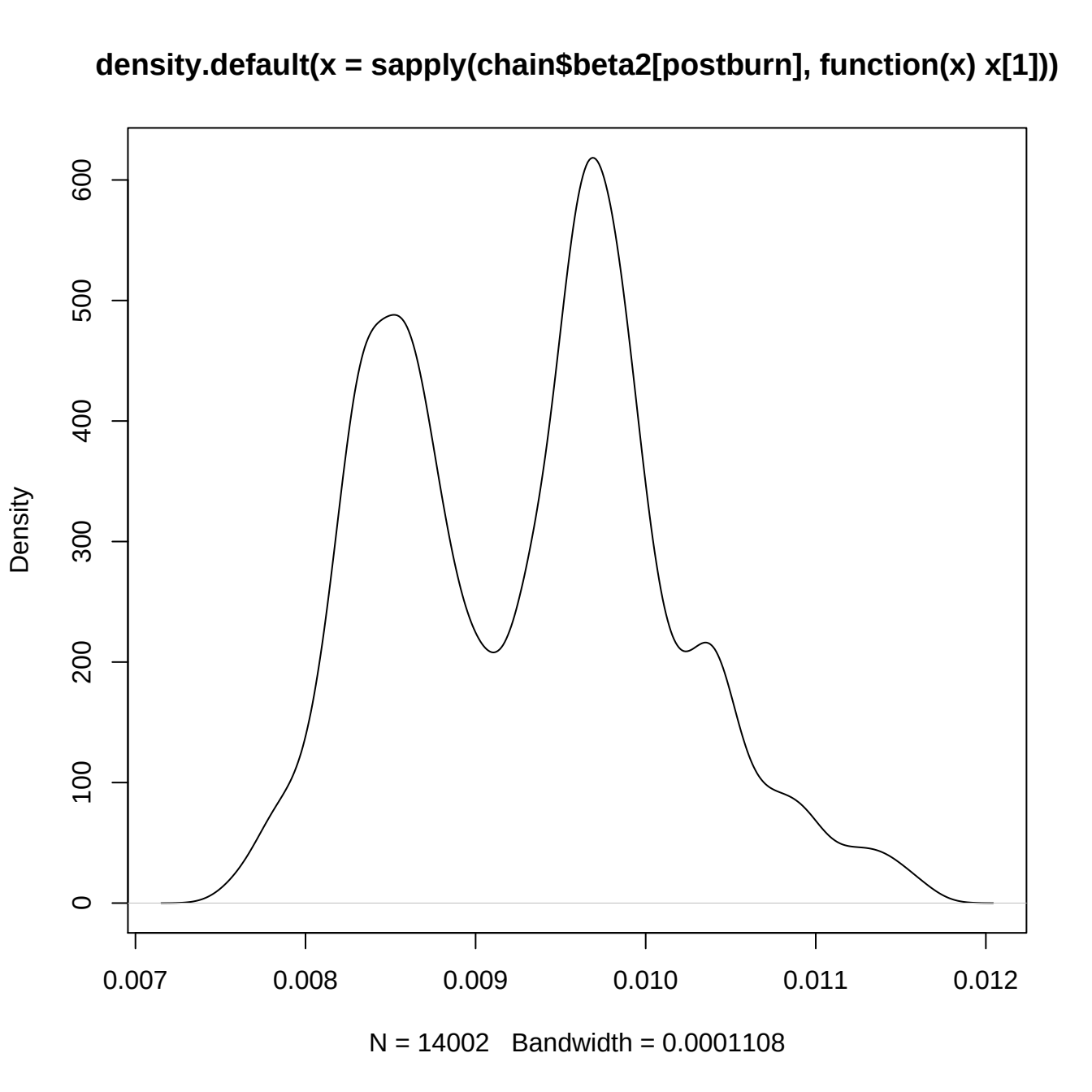
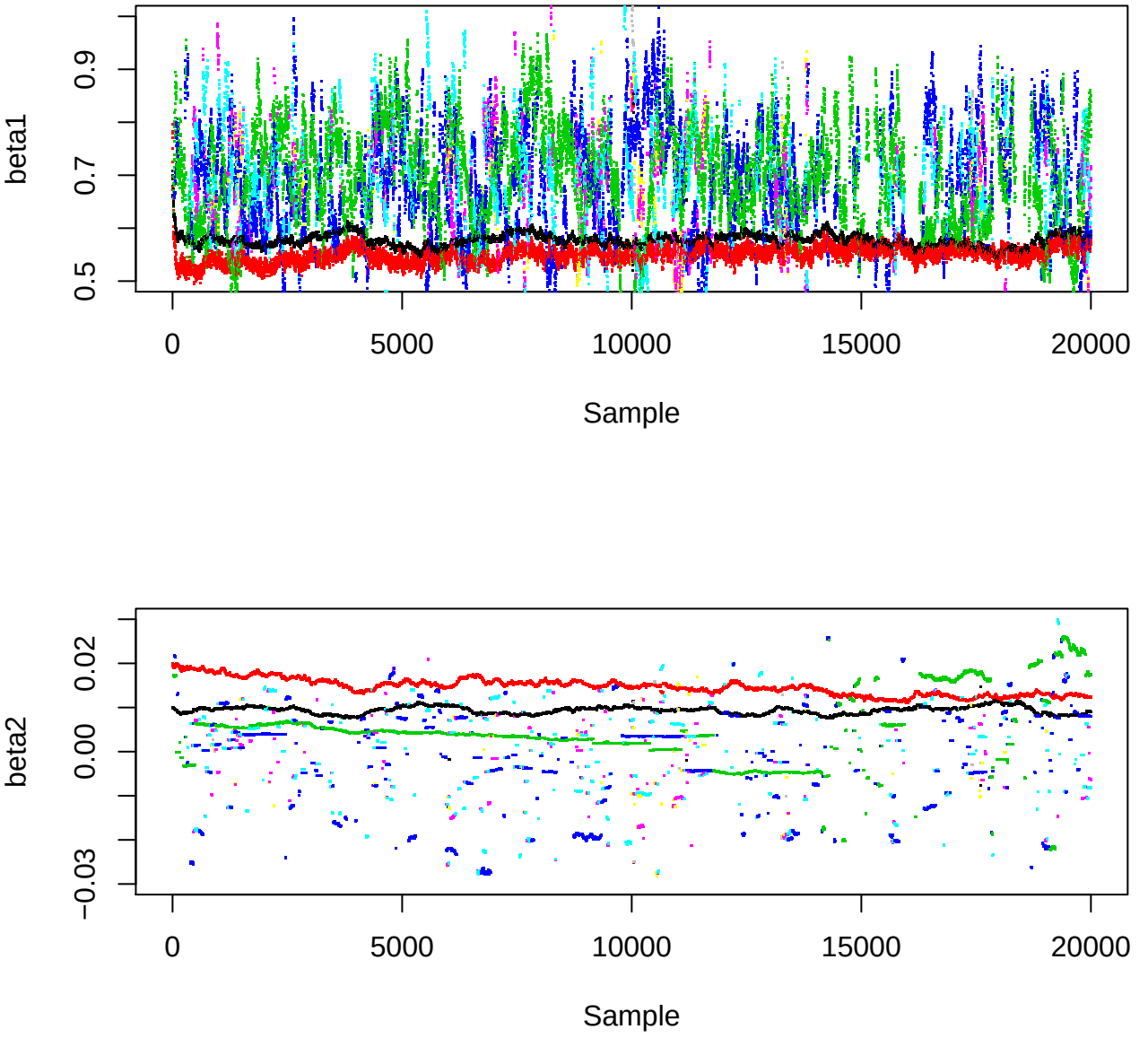
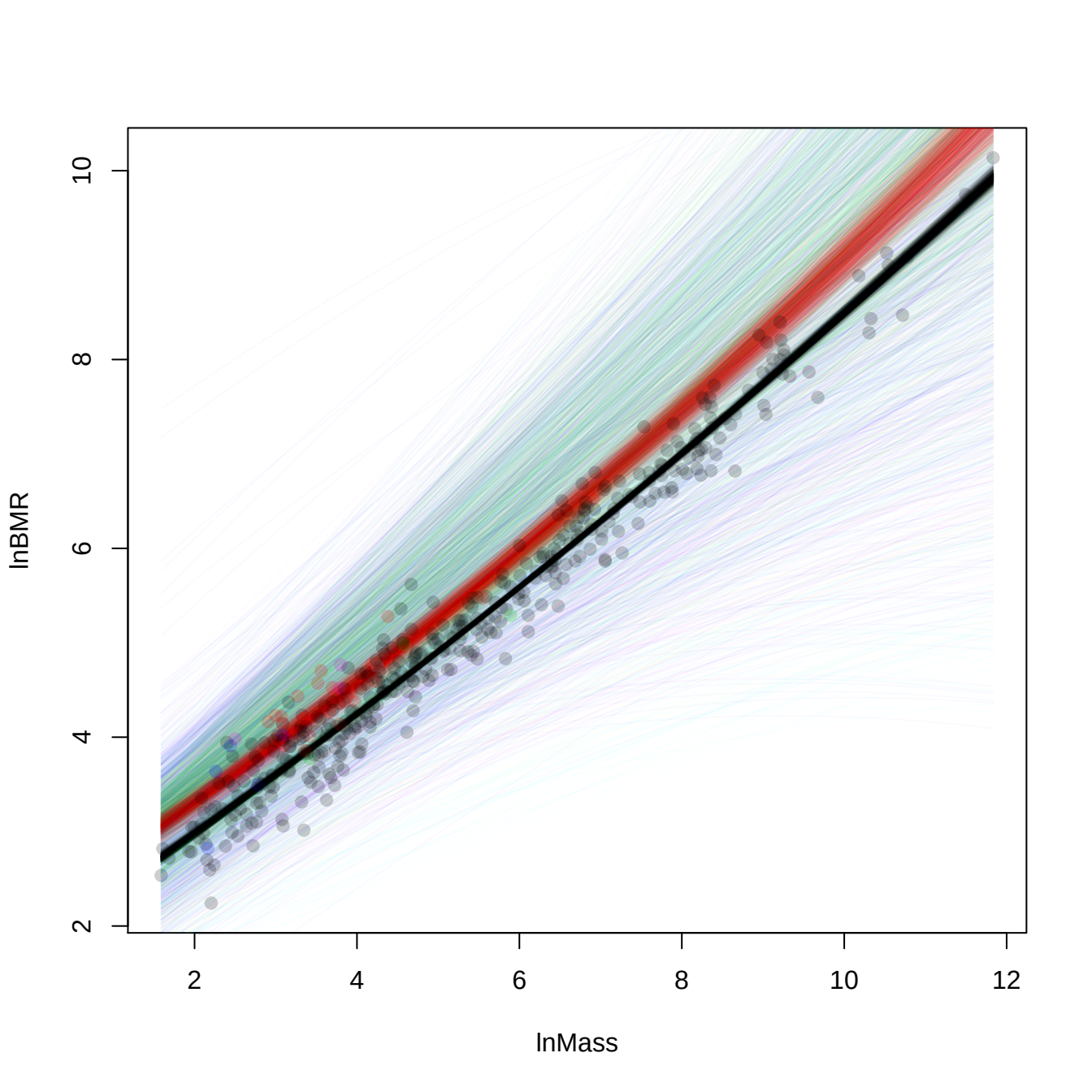

0.87
0.44
0
−0.44
−0.87


density.default(x = sapply(chain$beta2[postburn], function(x) x[1]))
Density
0
100
200
300
400
500
600
0.007
0.008
0.009
0.010
0.011
0.012
N = 14002 Bandwidth = 0.0001108


beta1
0.5
0.7
0.9
0
5000
10000
15000
20000
Sample
beta2
−0.03
0.00
0.02
0
5000
10000
15000
20000
Sample


lnBMR
lnMass
2
4
6
8
10
12