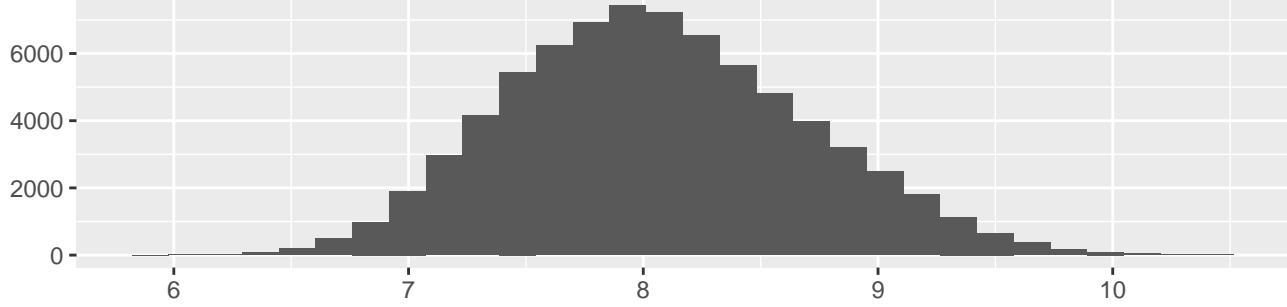
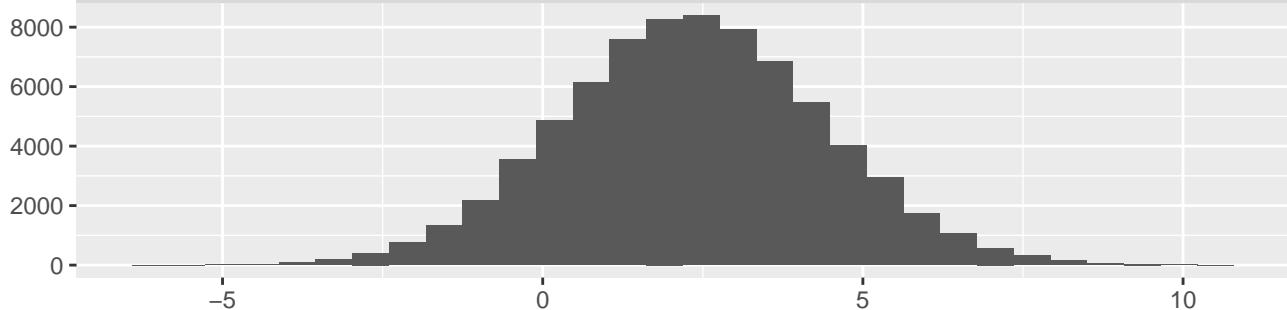


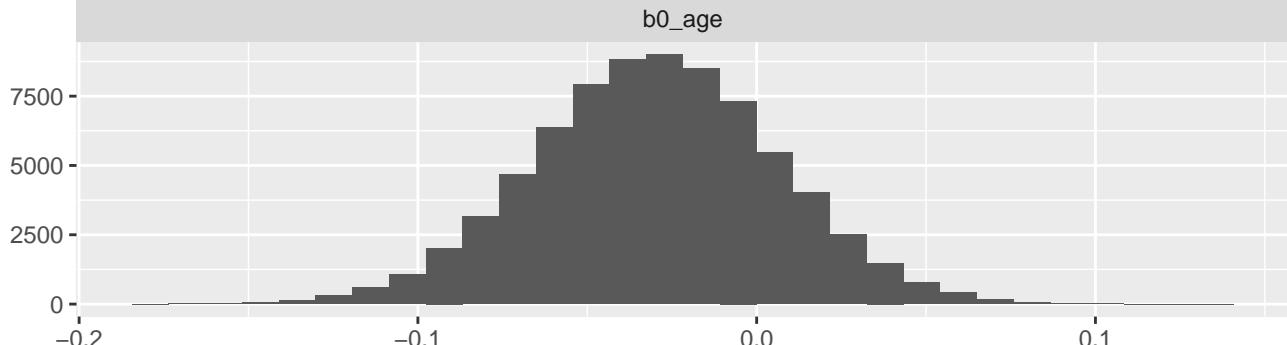
b.rand.sigma



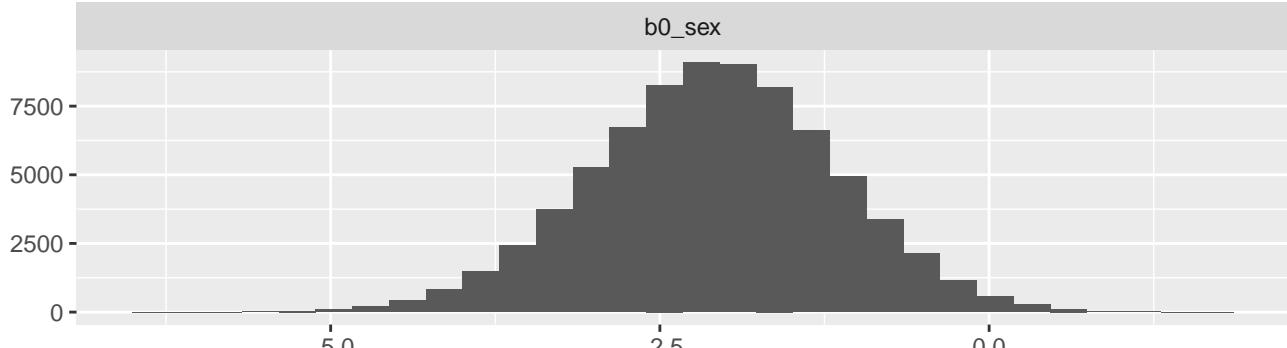
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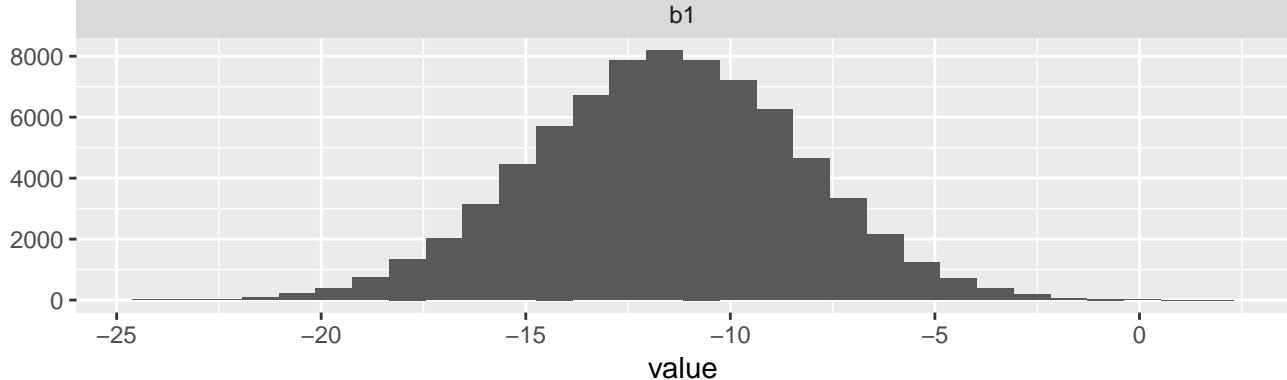
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b0_sex

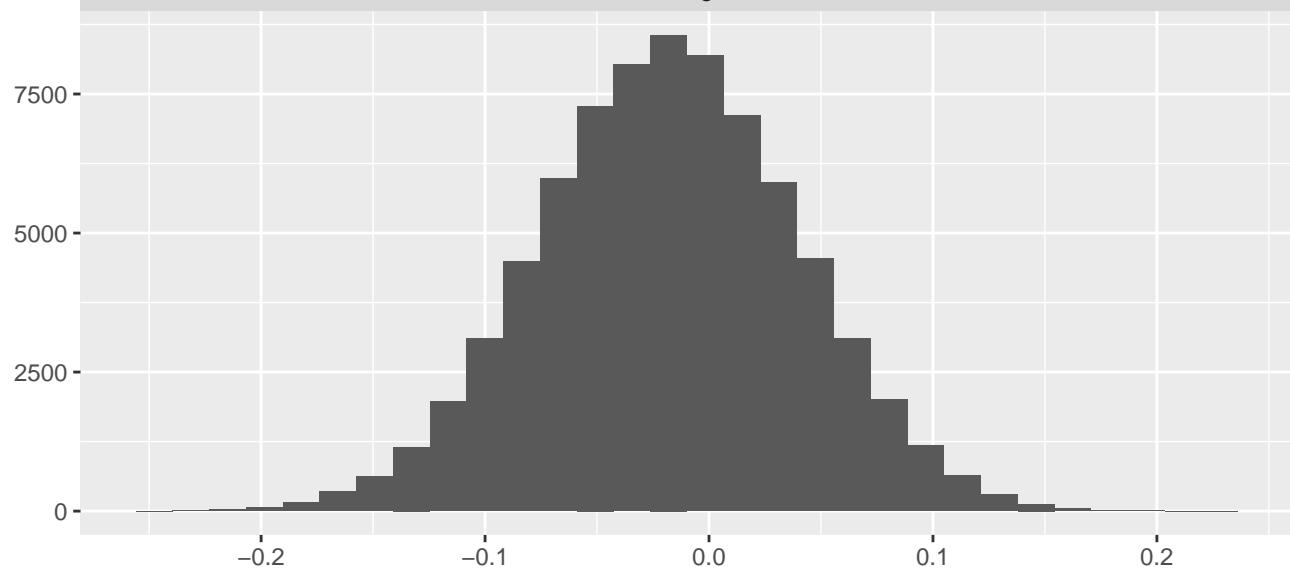


b1

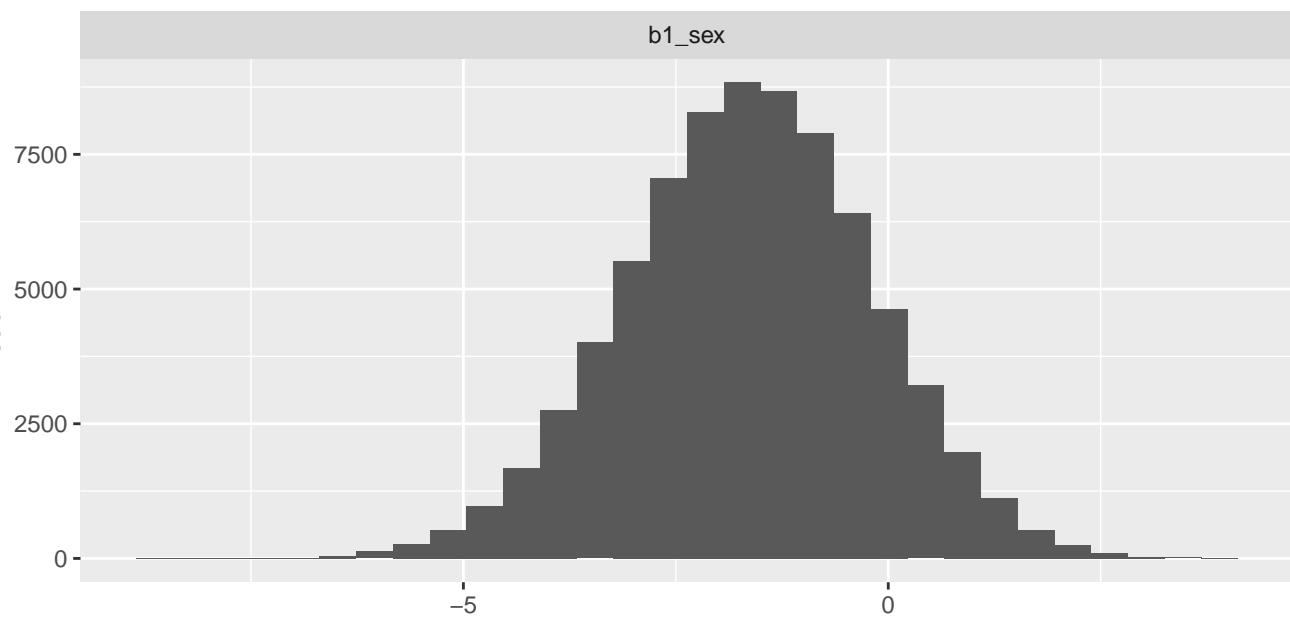


value

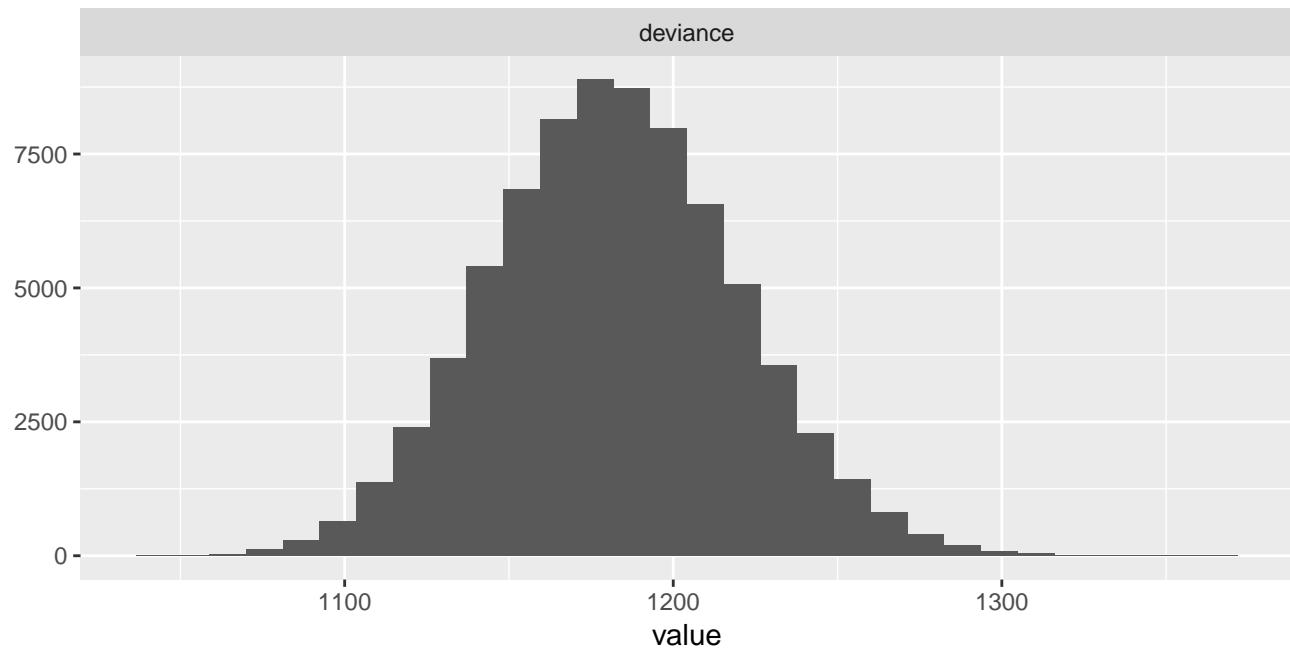
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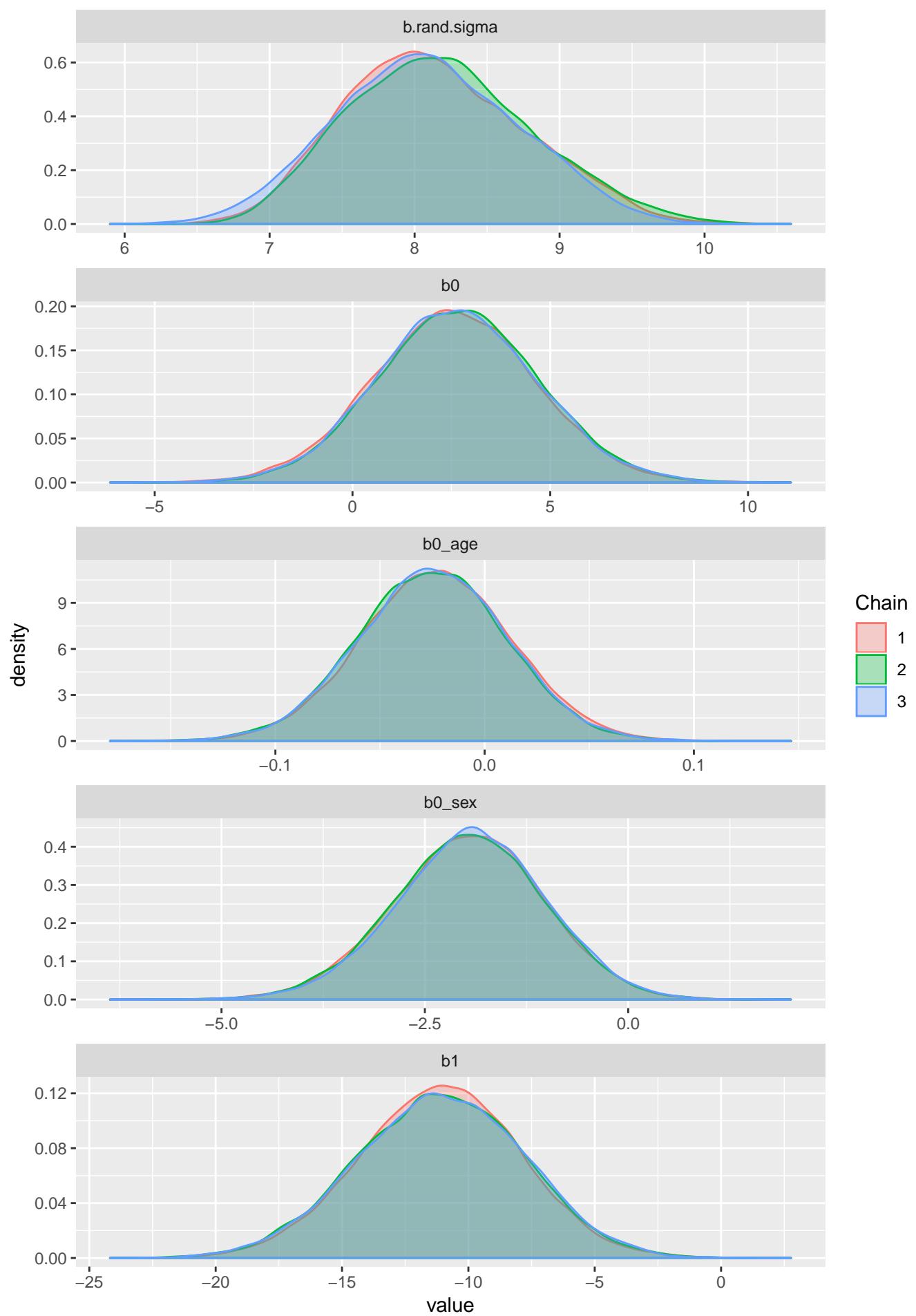


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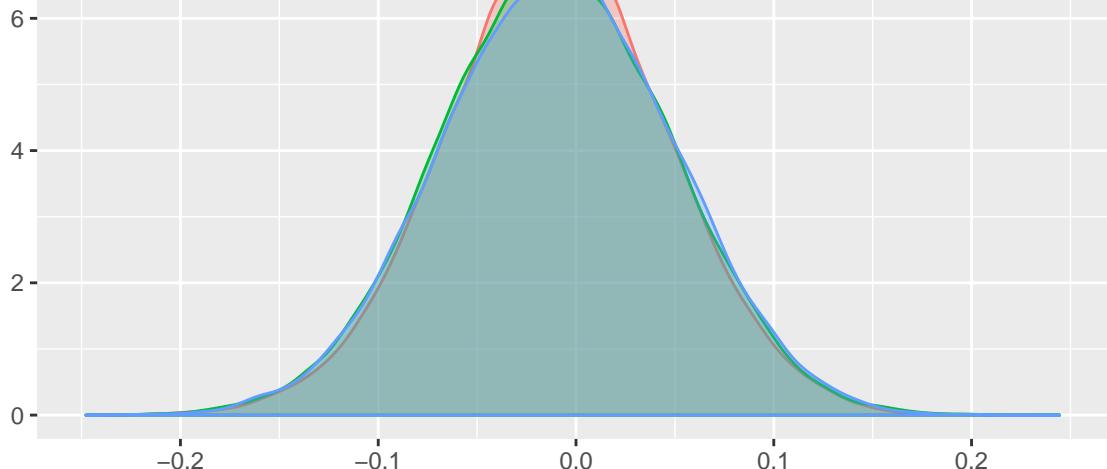


deviance





b1_age

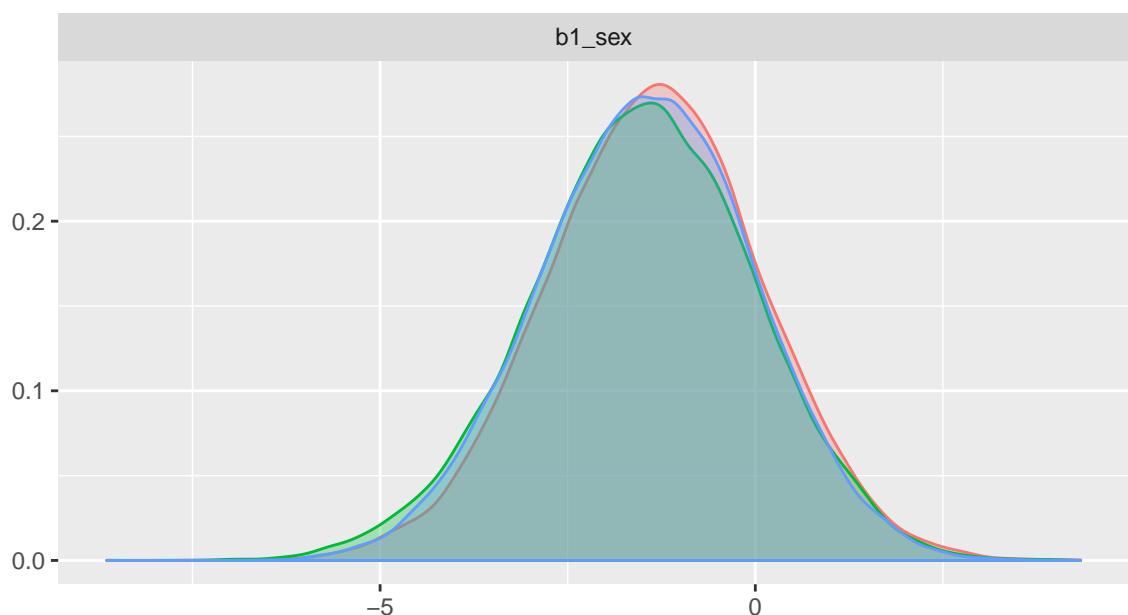


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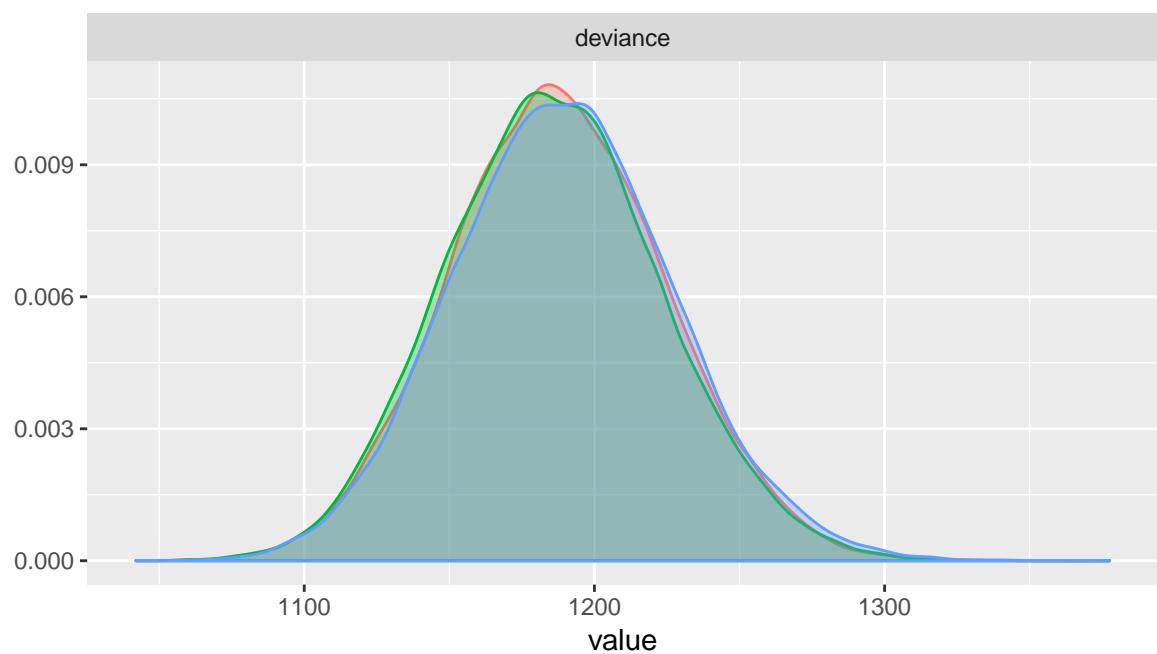
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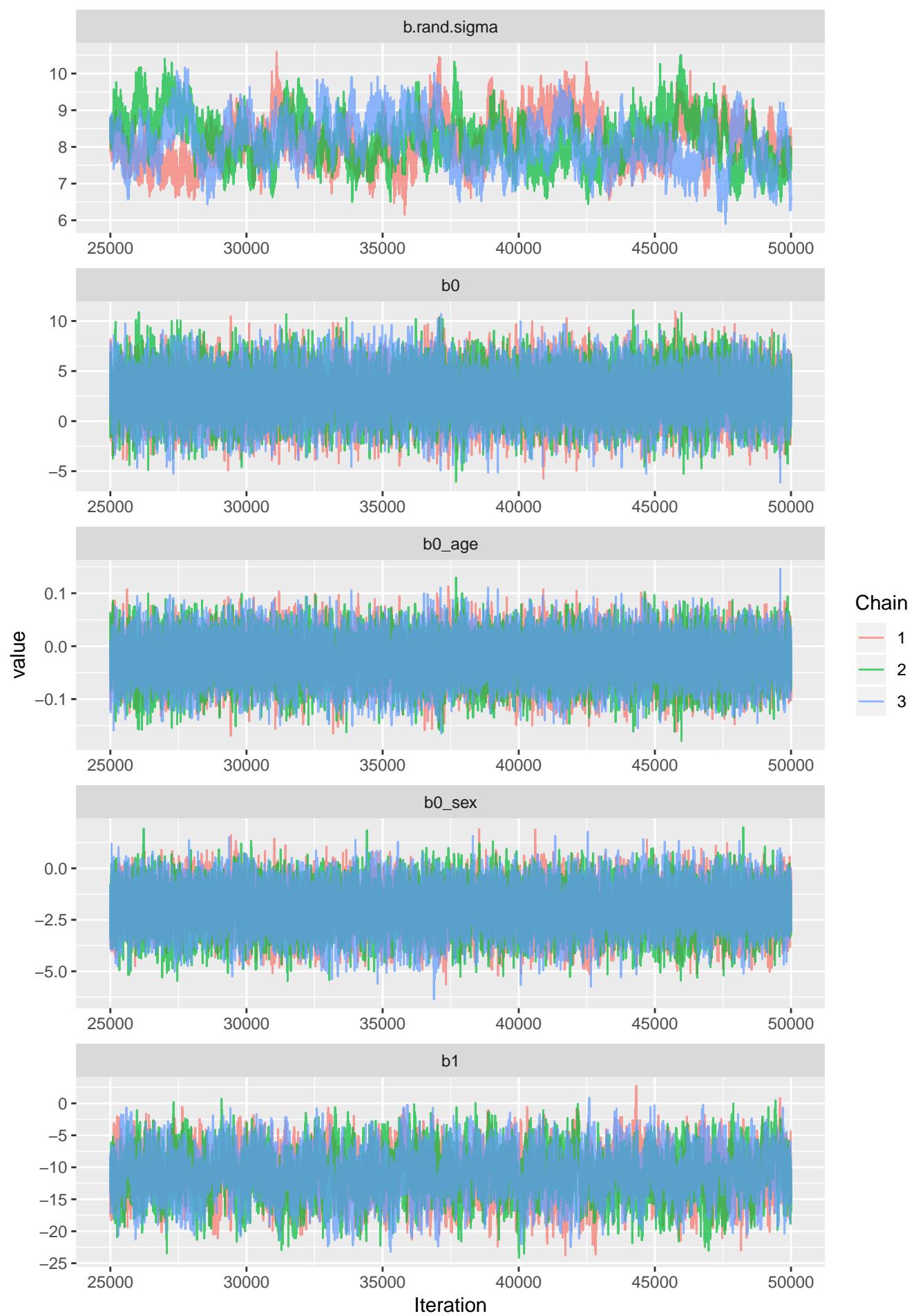
Chain

- 1
- 2
- 3

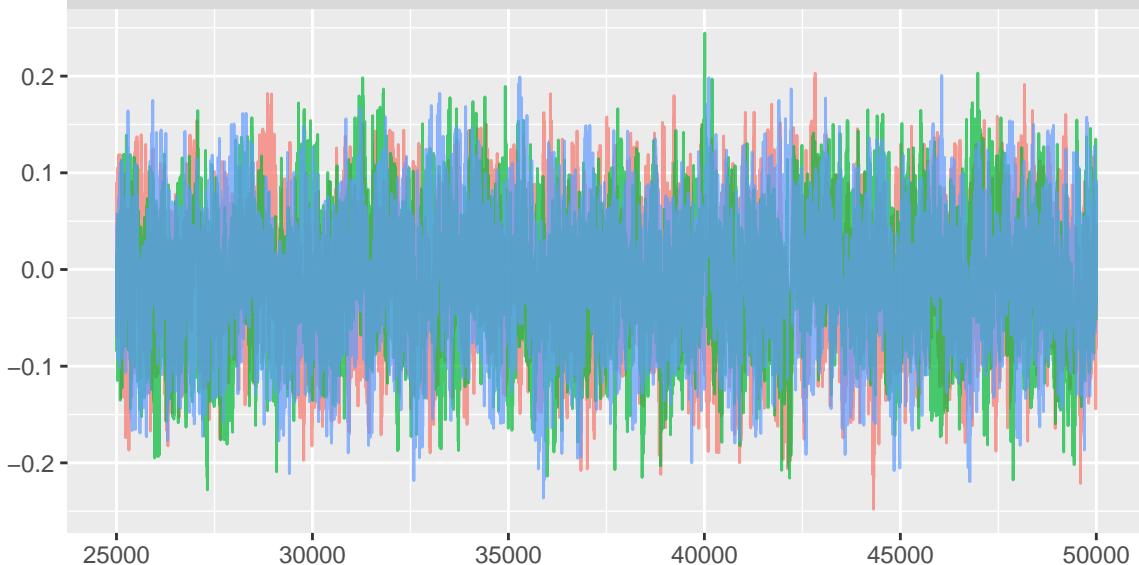


deviance

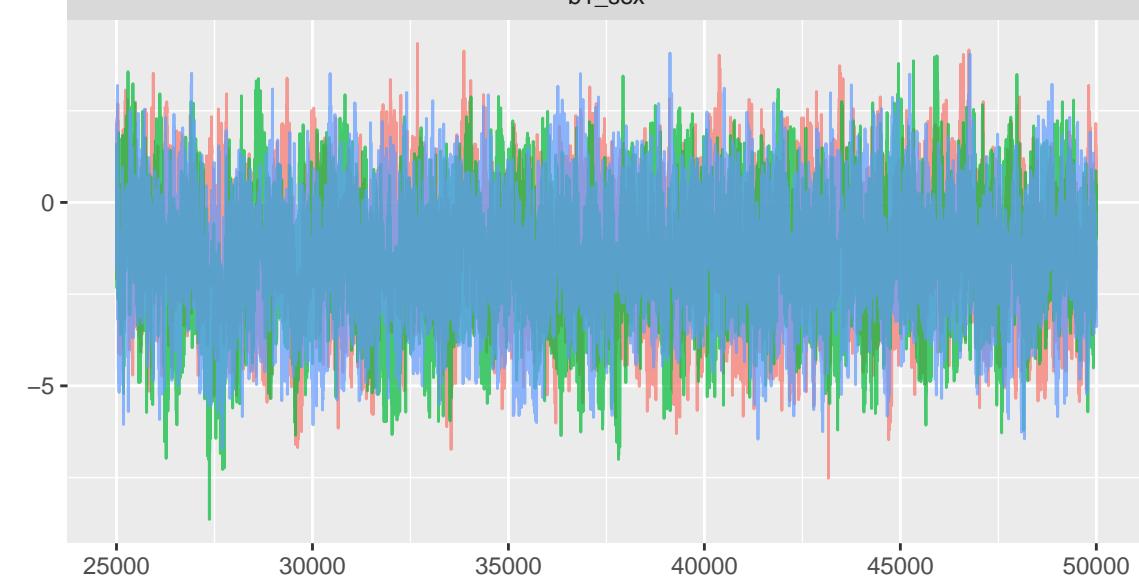




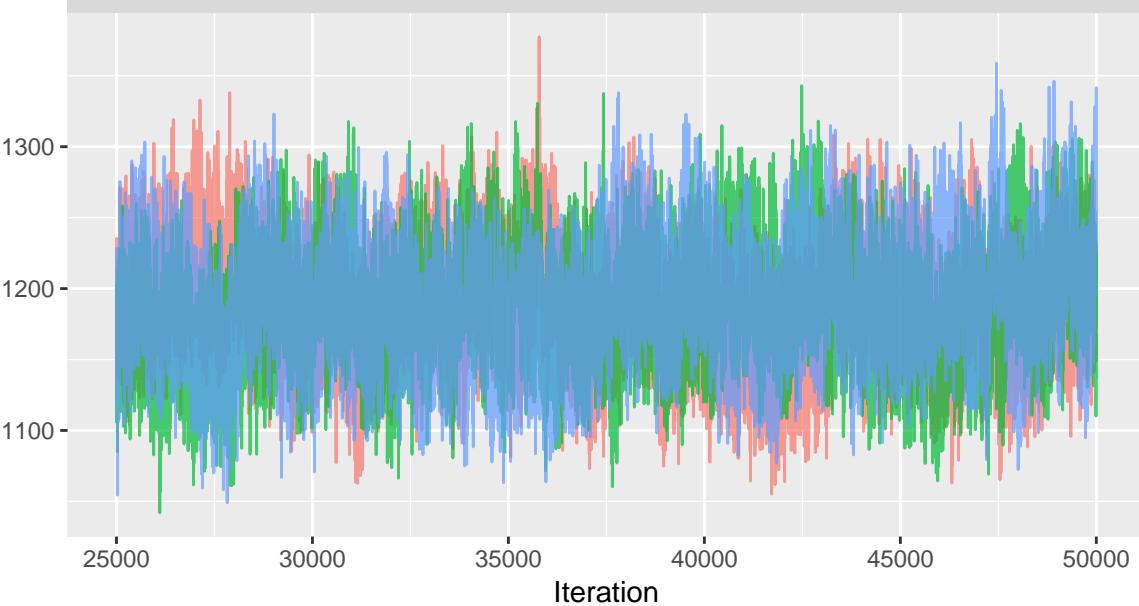
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b1_sex

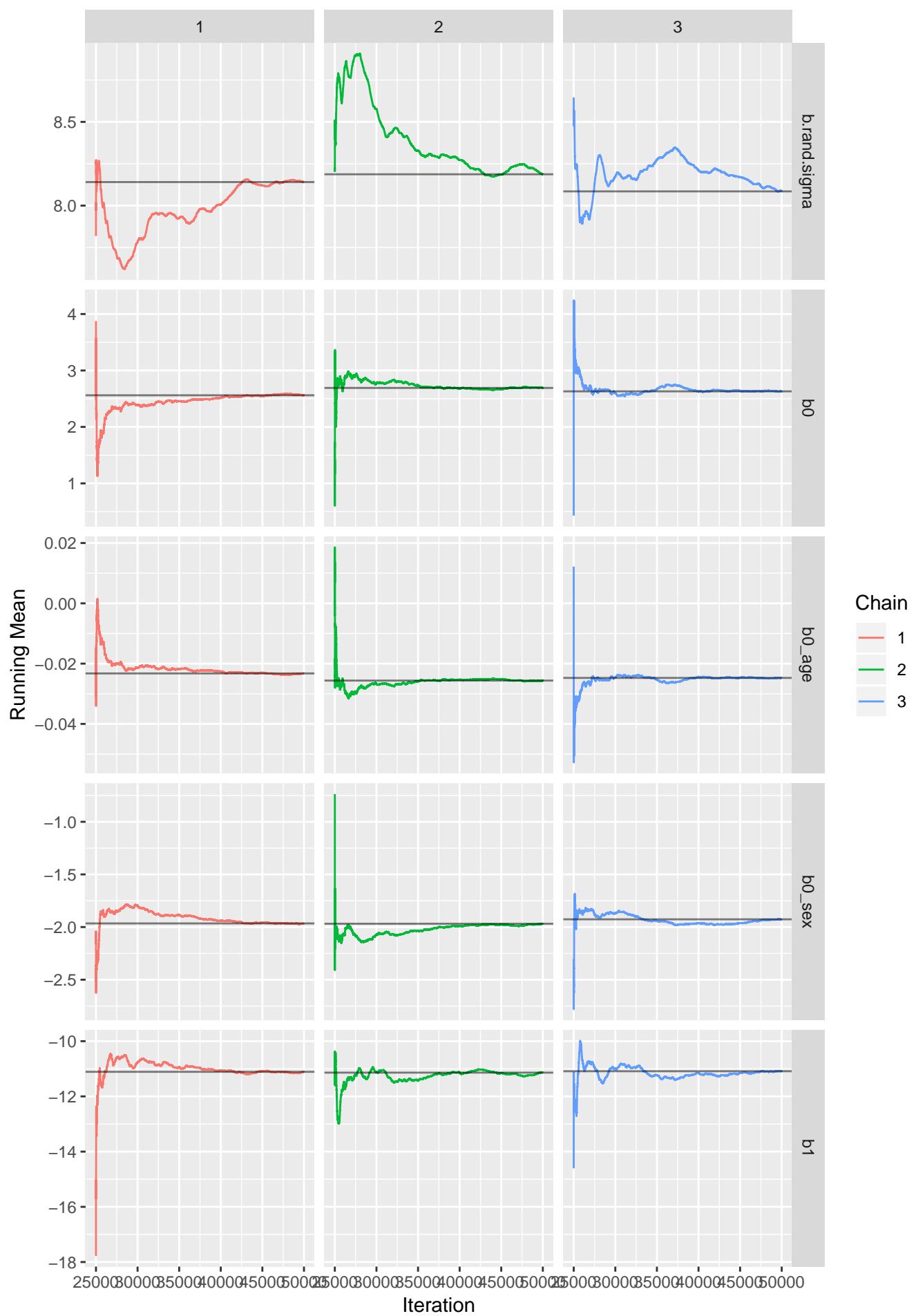


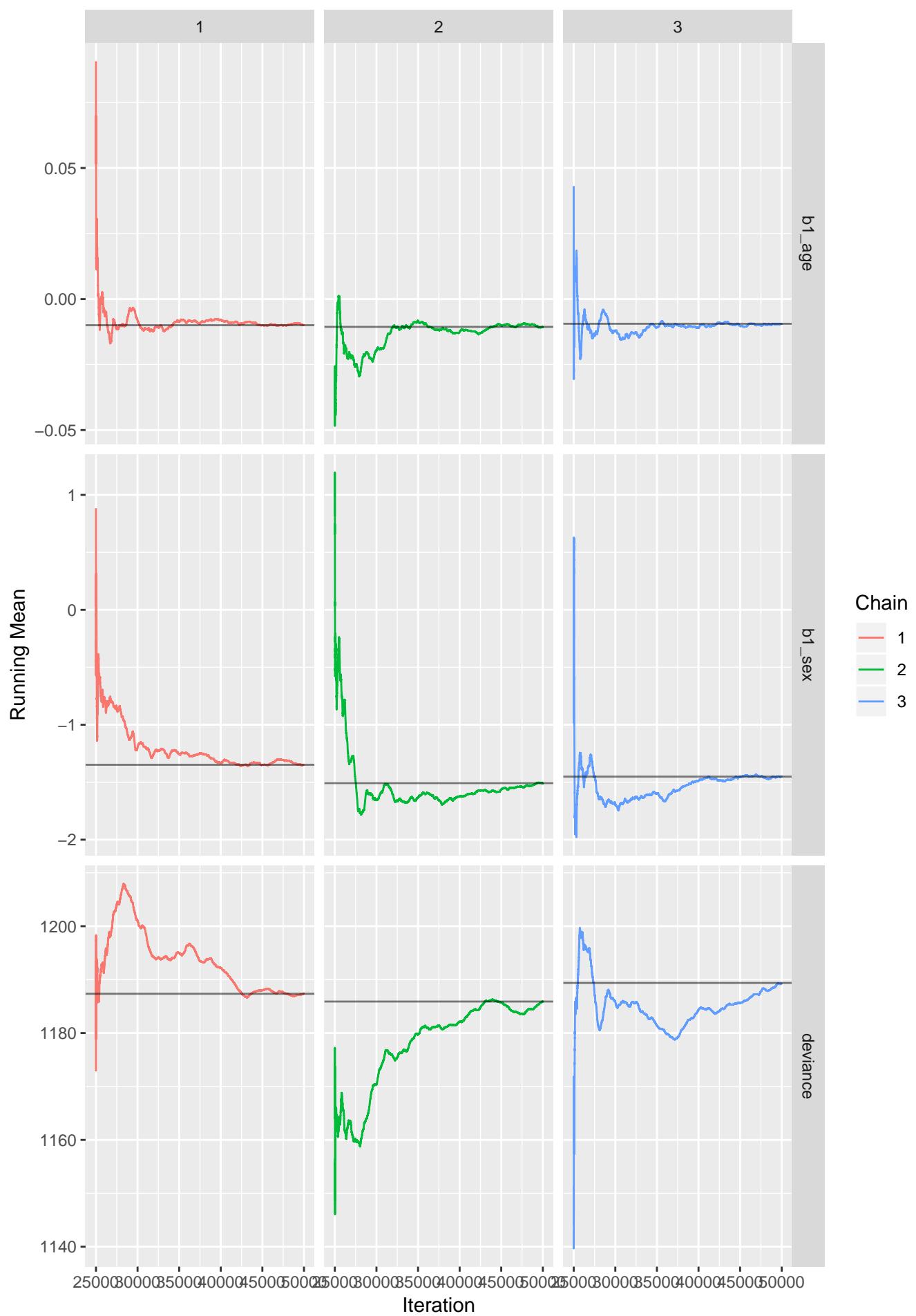
deviance

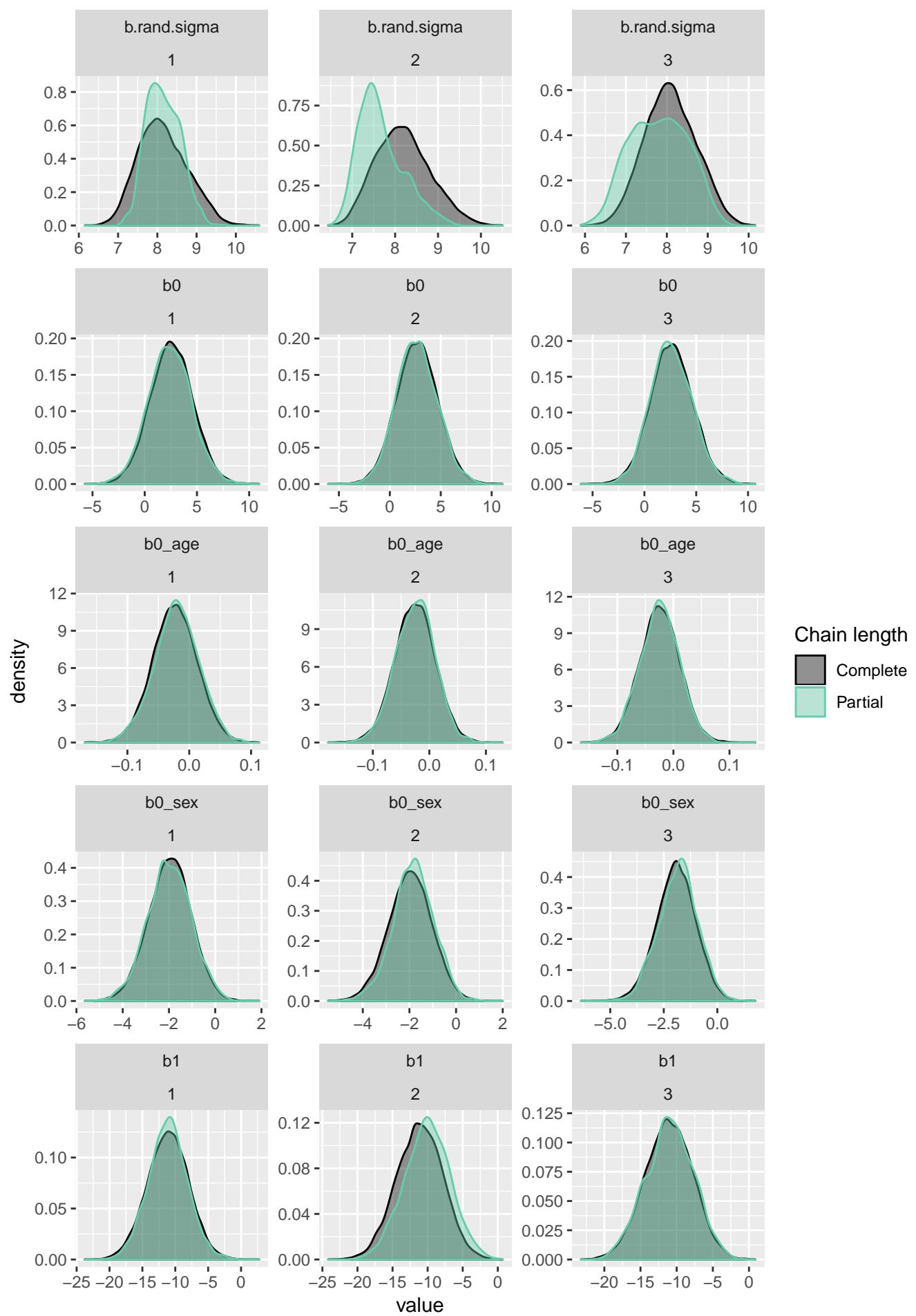


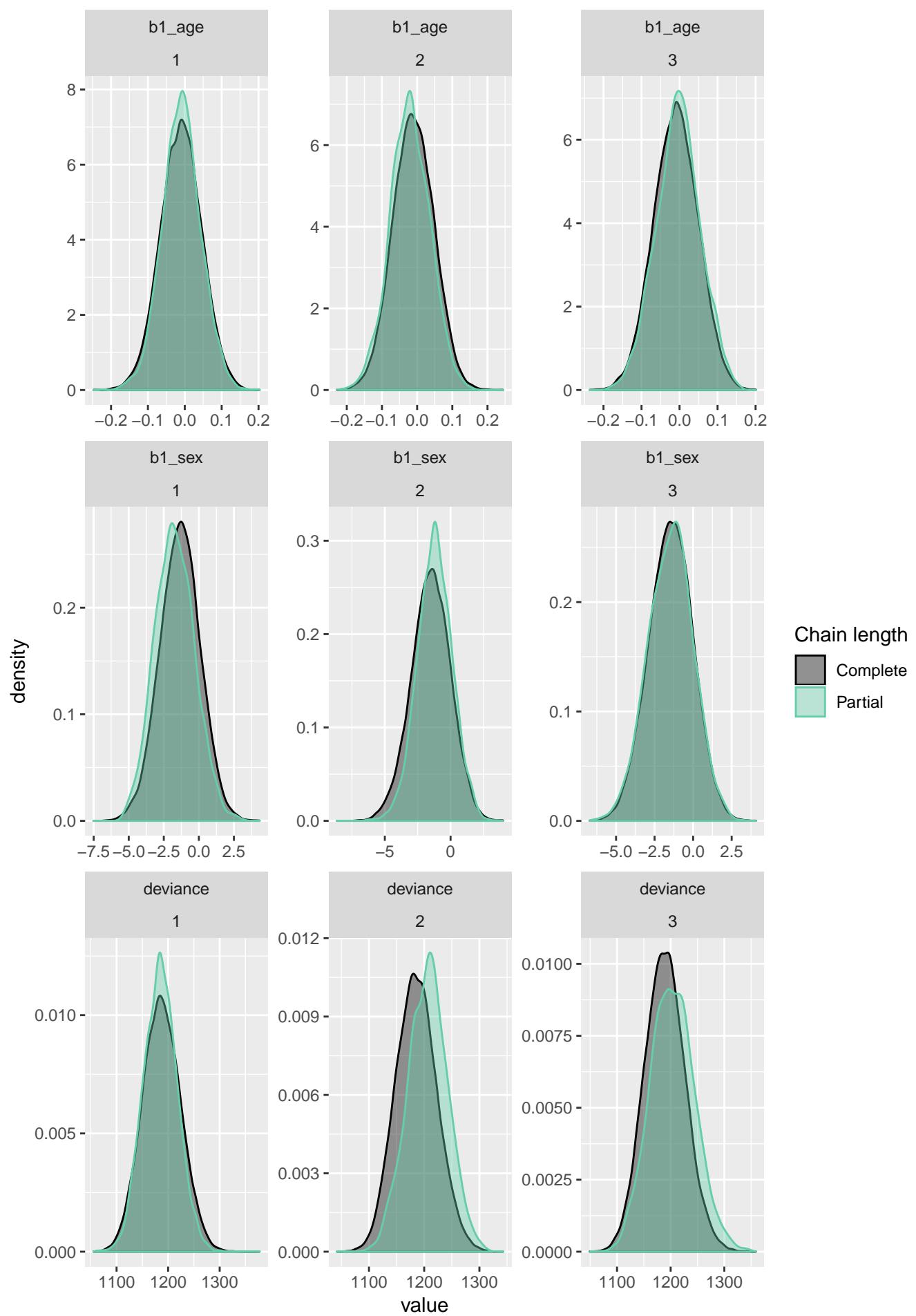
Chain
1
2
3

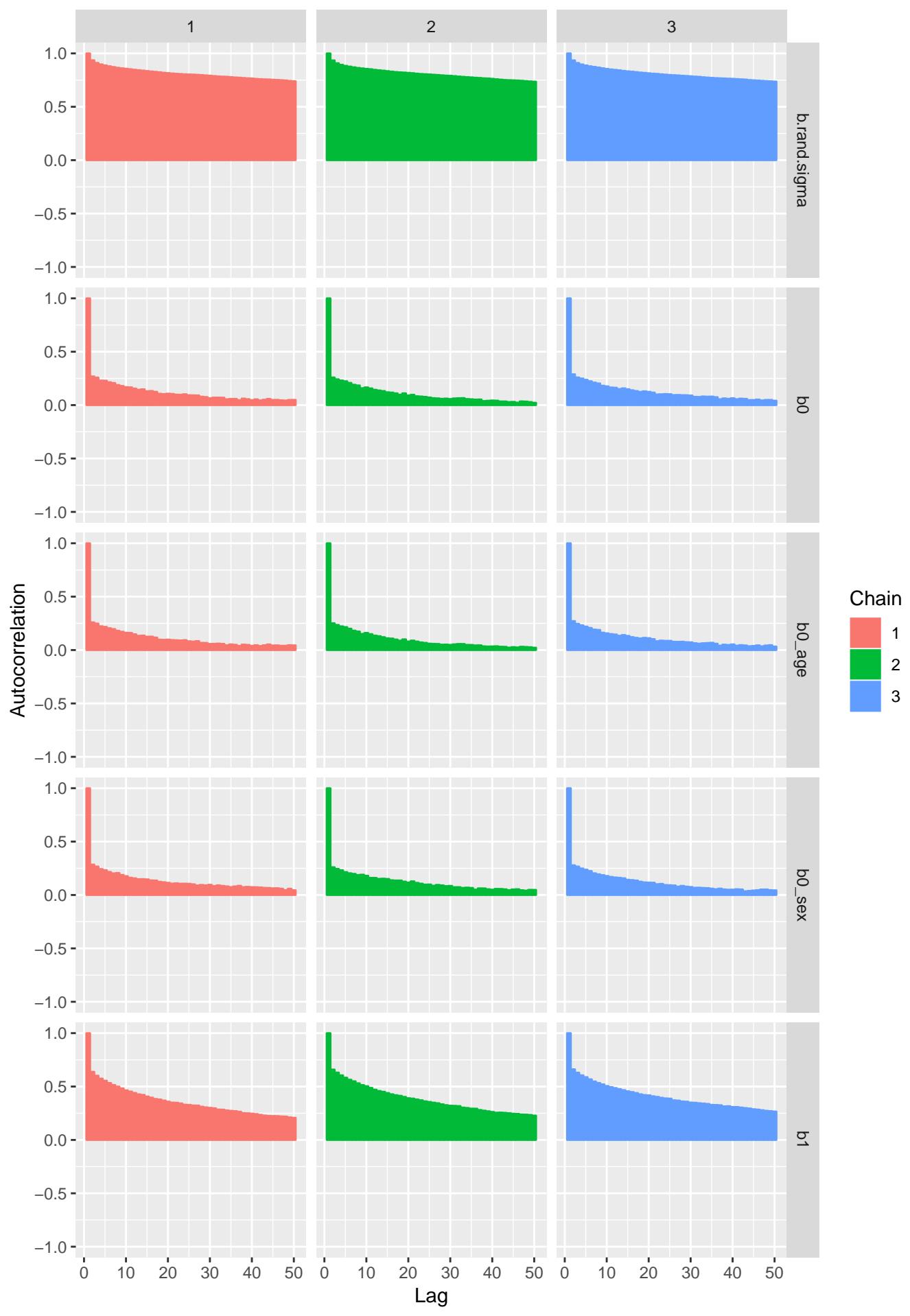
Iteration

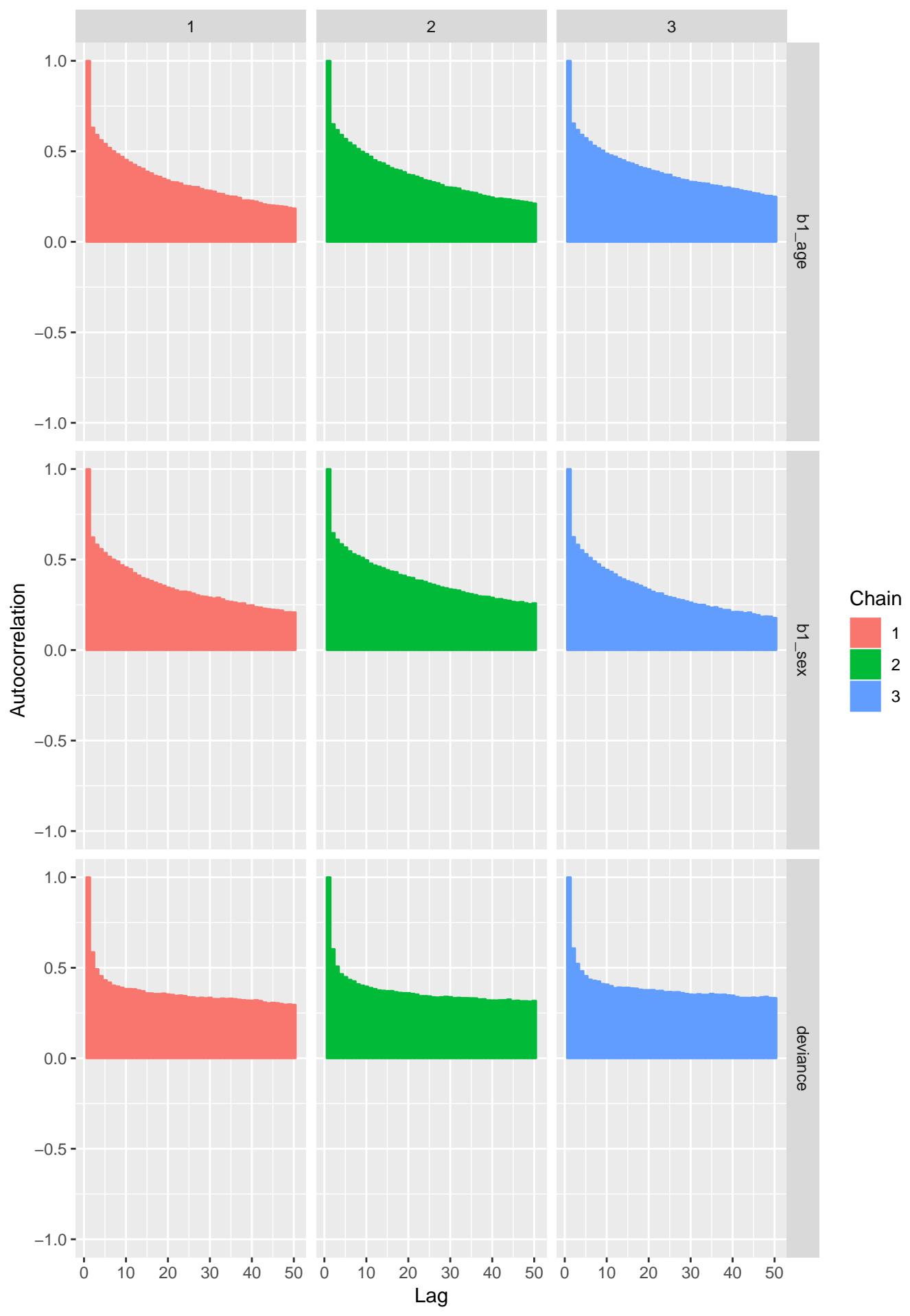


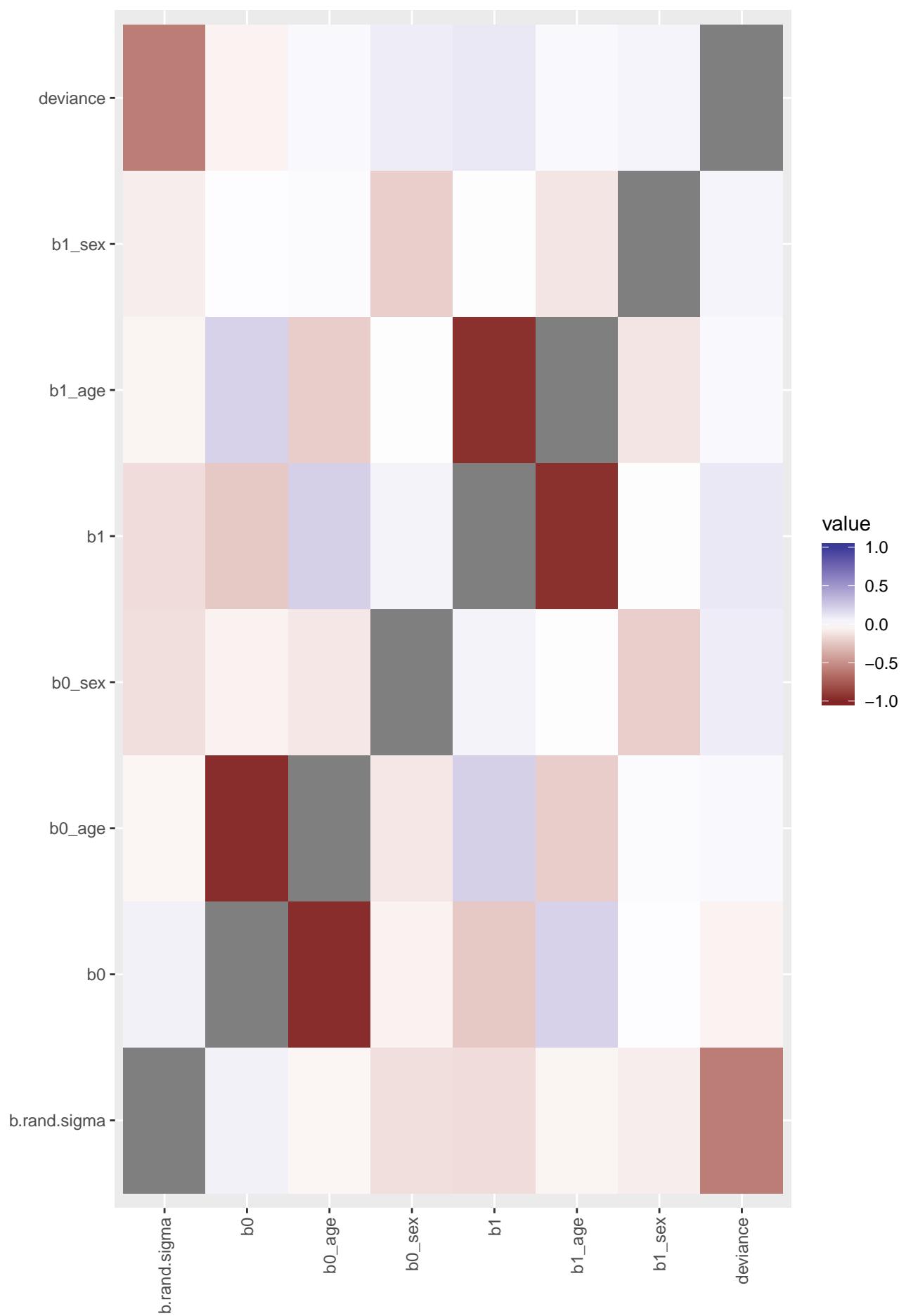




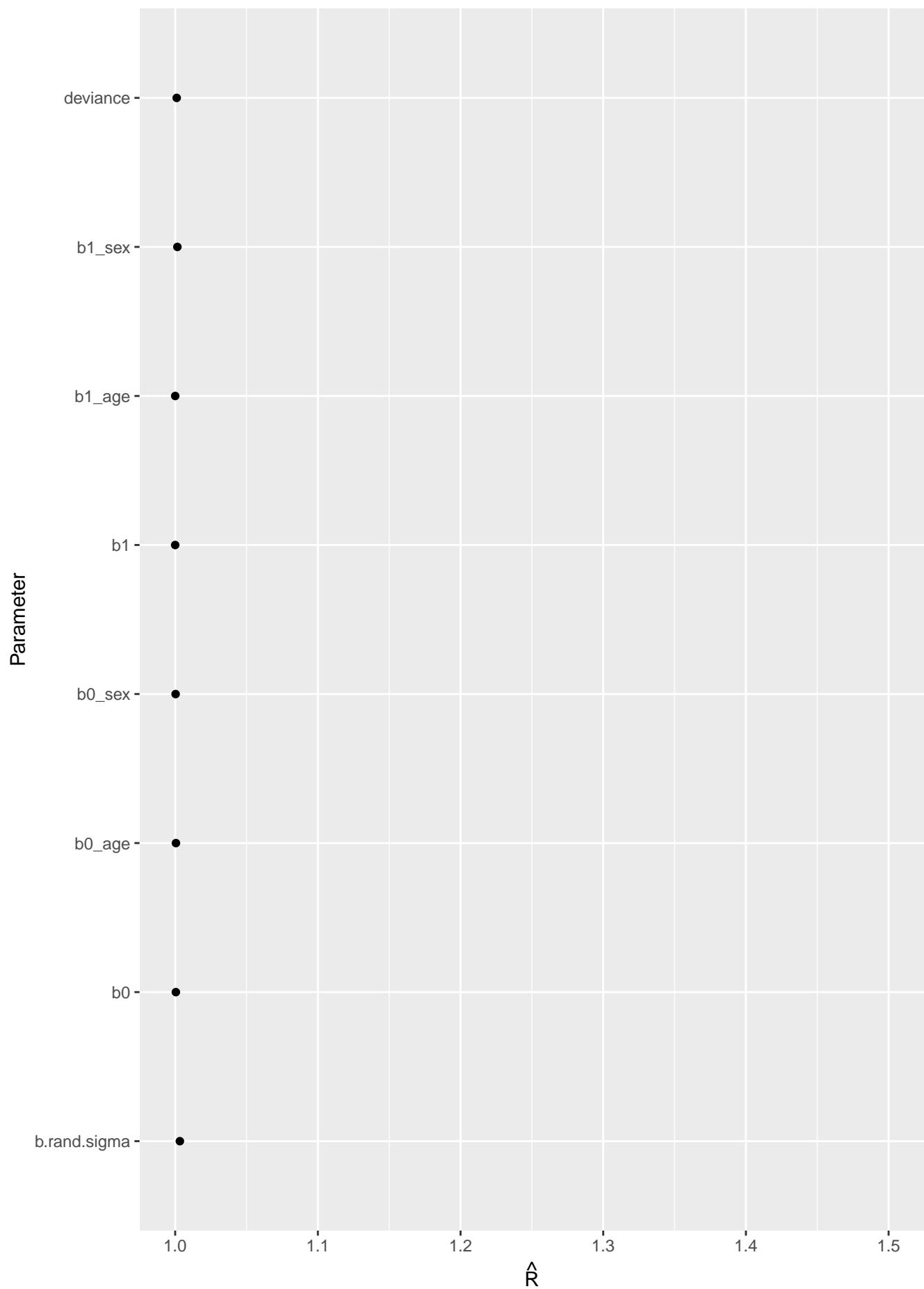








Potential Scale Reduction Factors



Geweke Diagnostics

