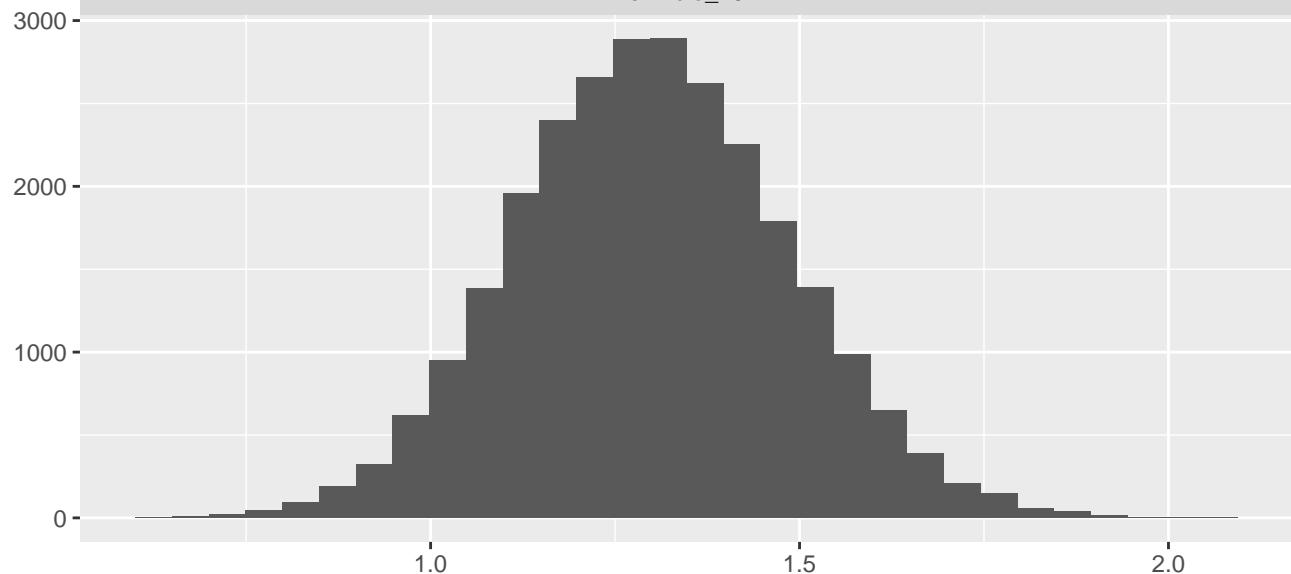
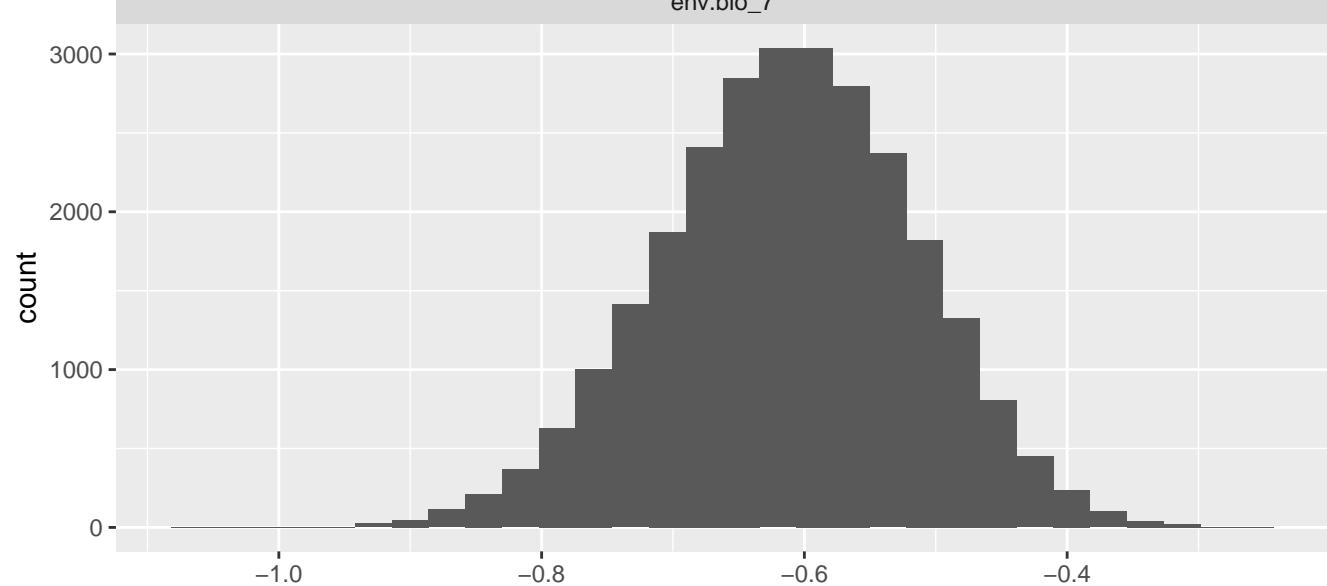


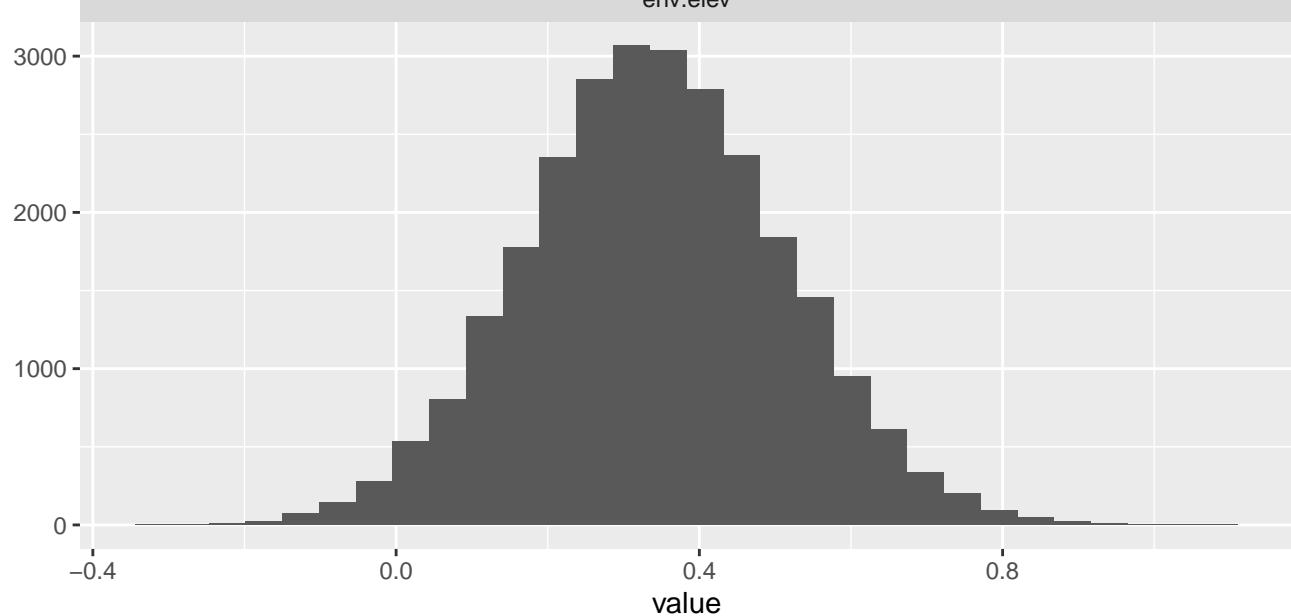
env.bio_15



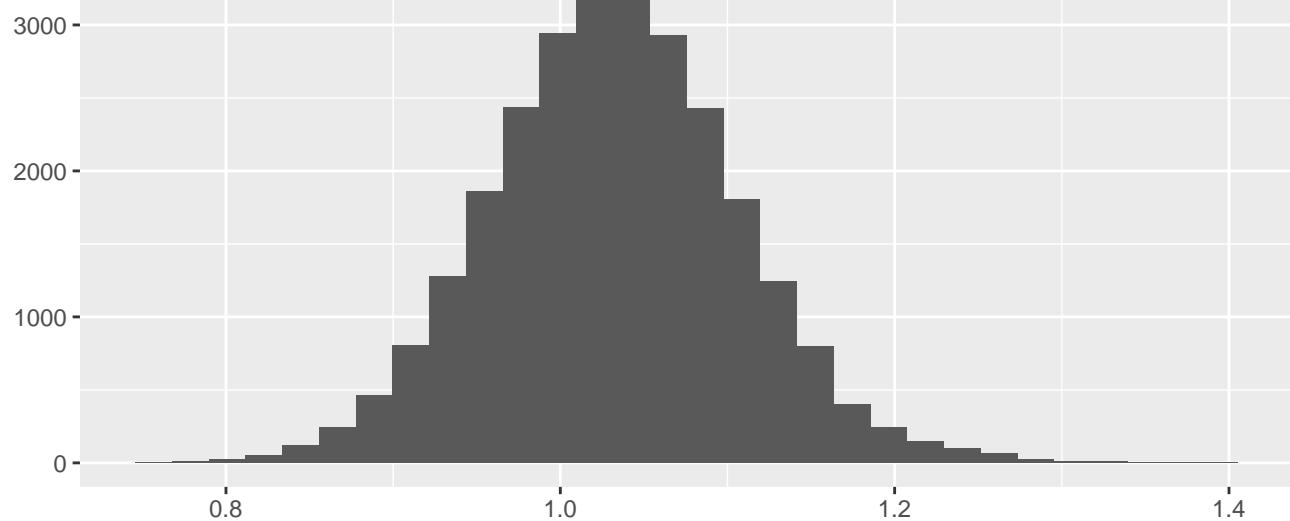
env.bio_7



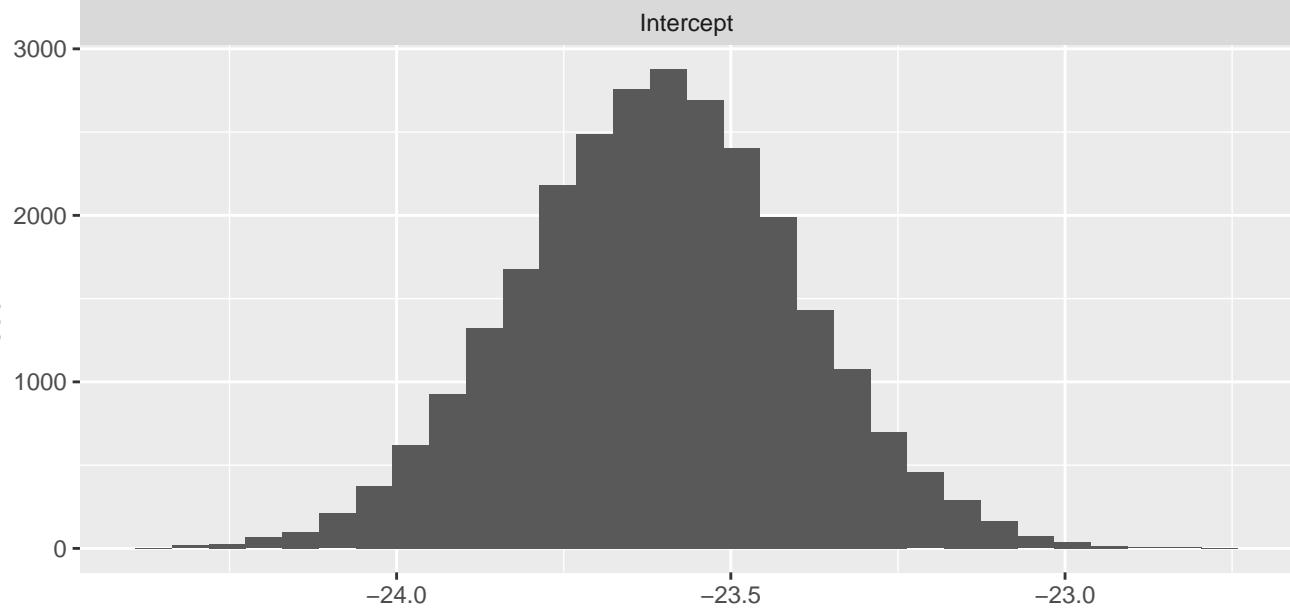
env.elev



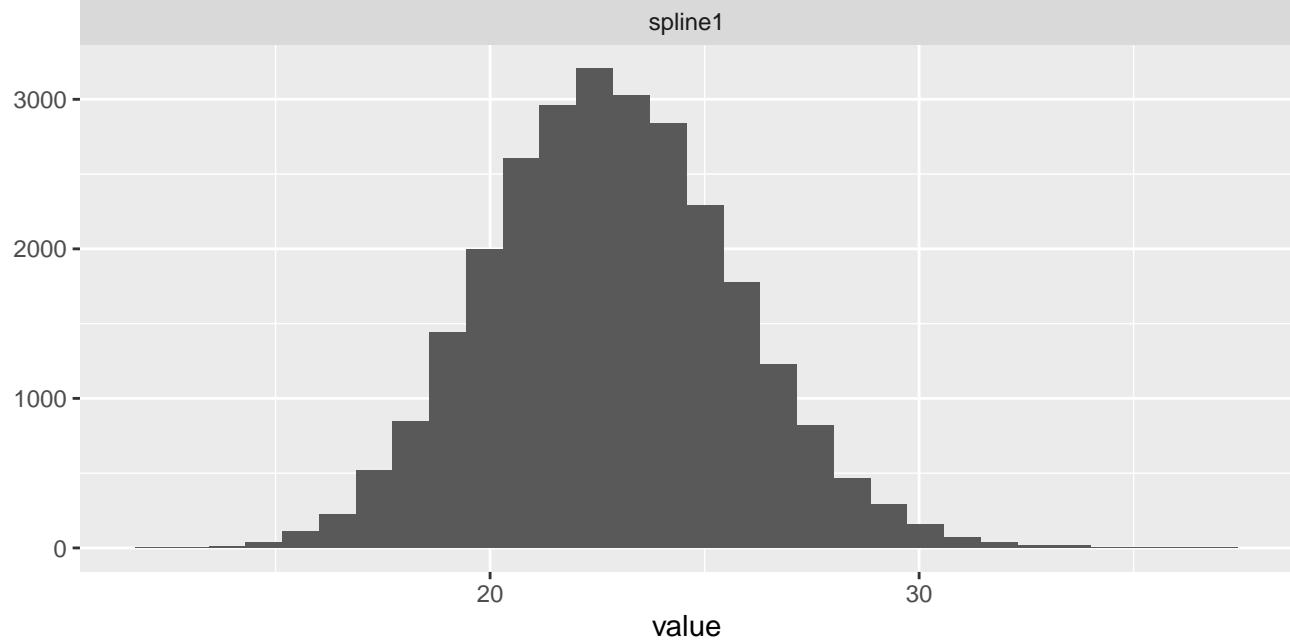
env.npp



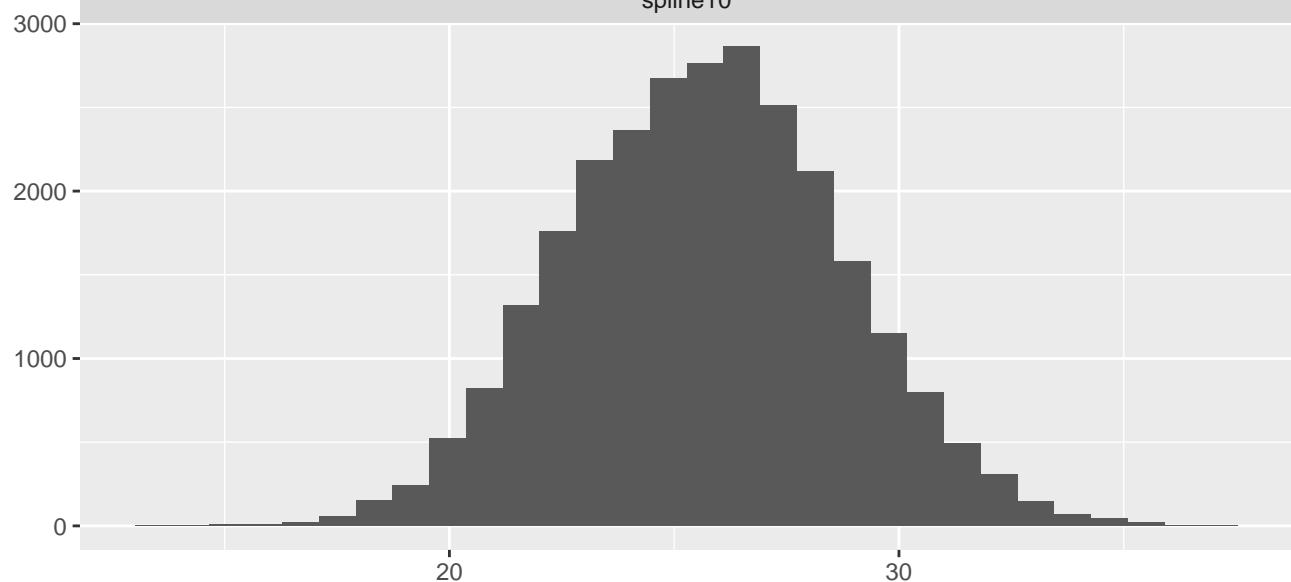
Intercept



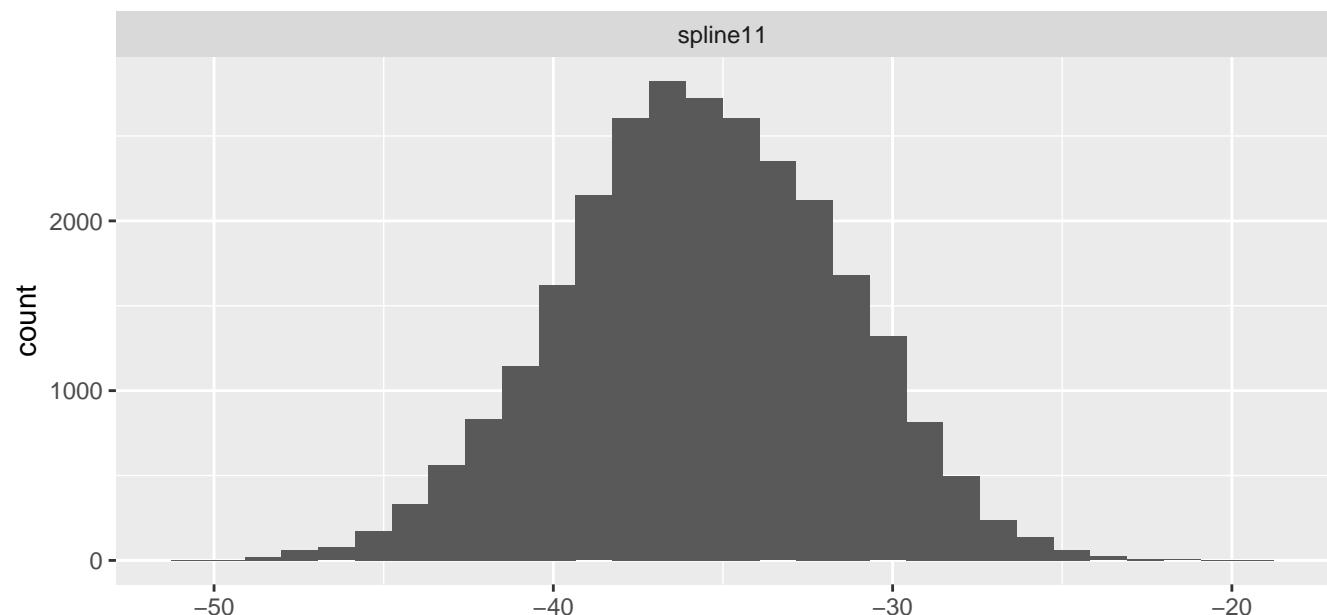
spline1



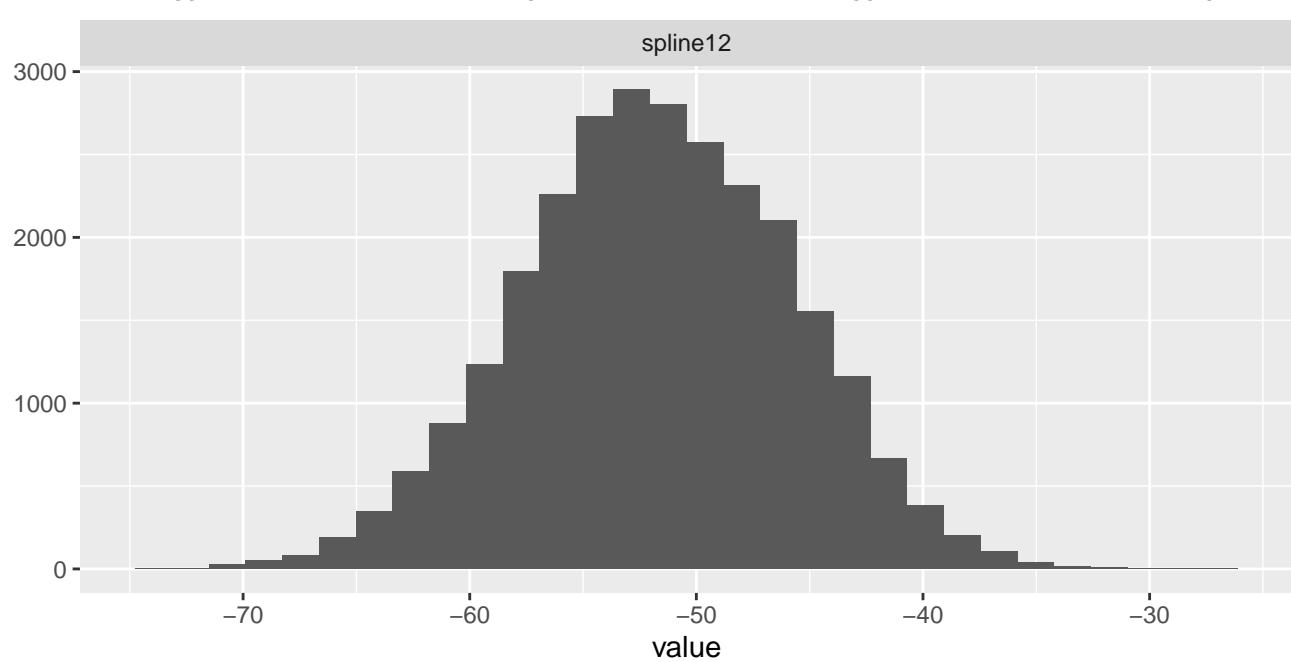
spline10



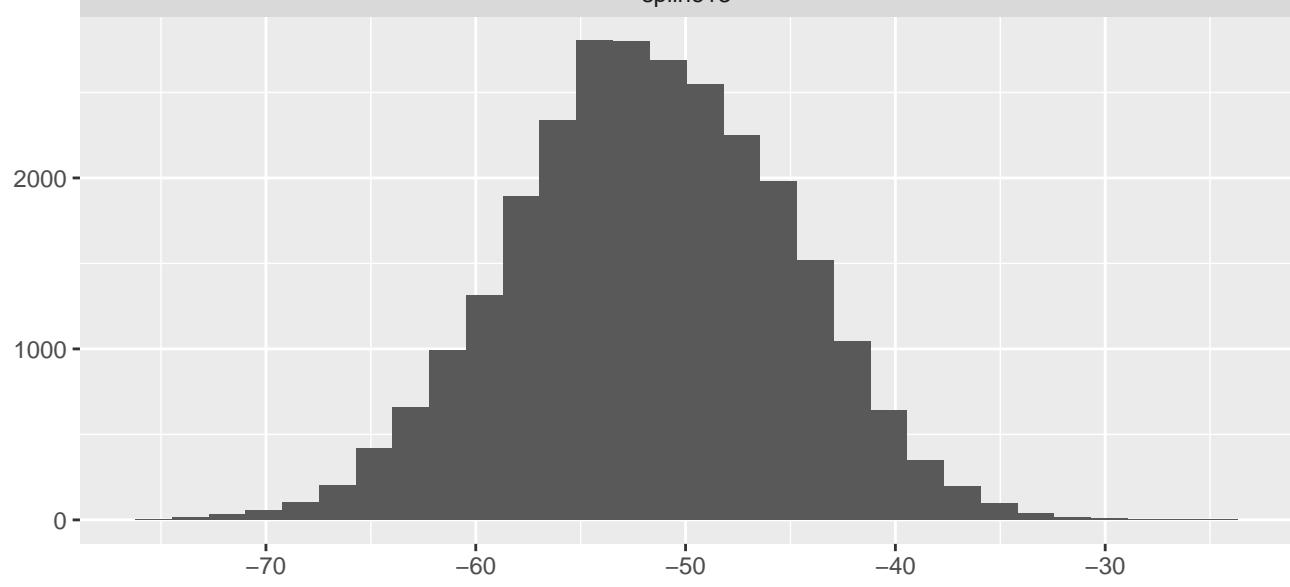
spline11



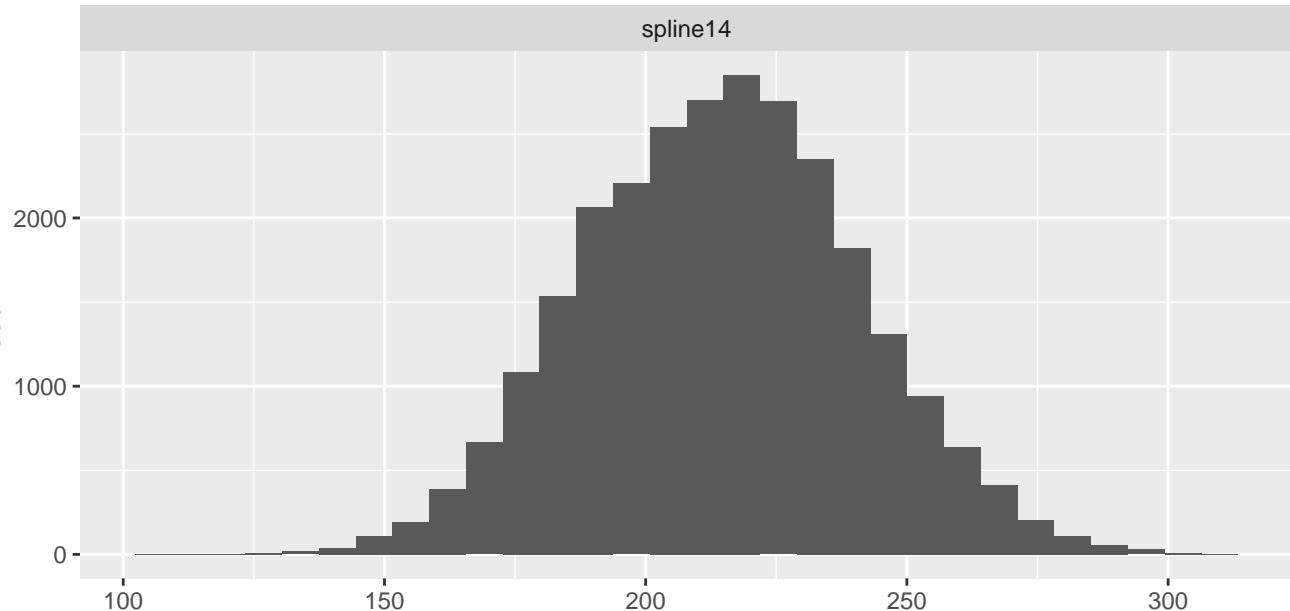
spline12



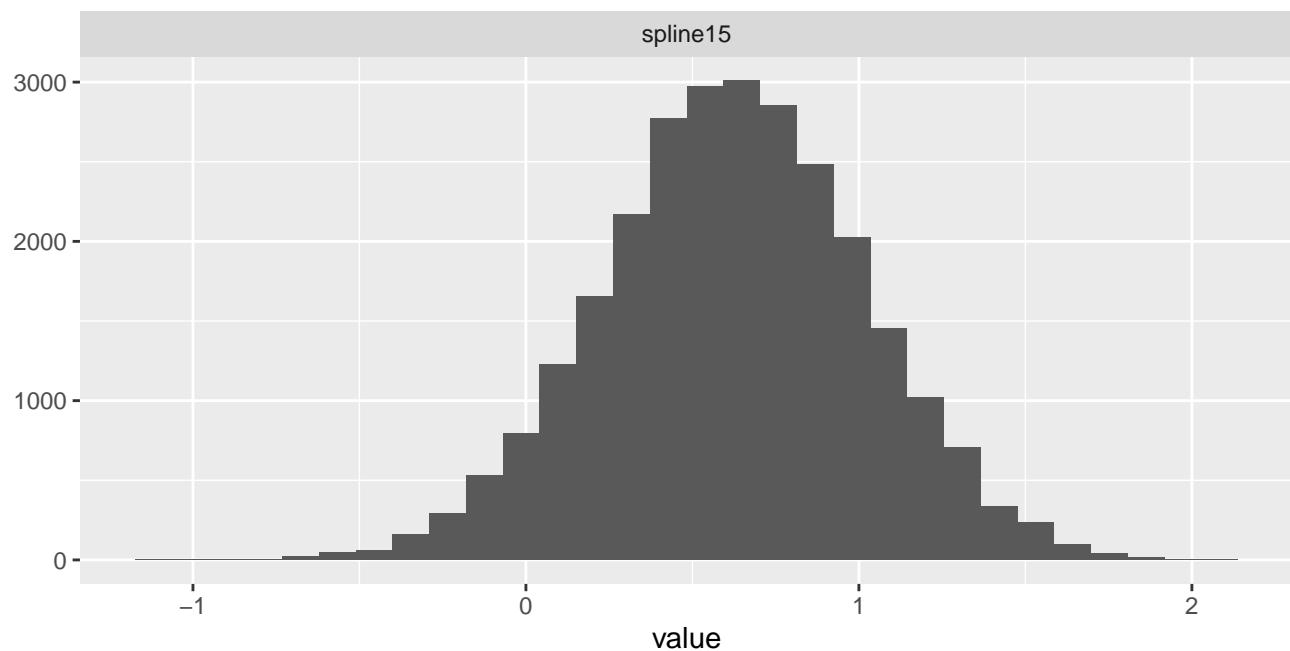
spline13



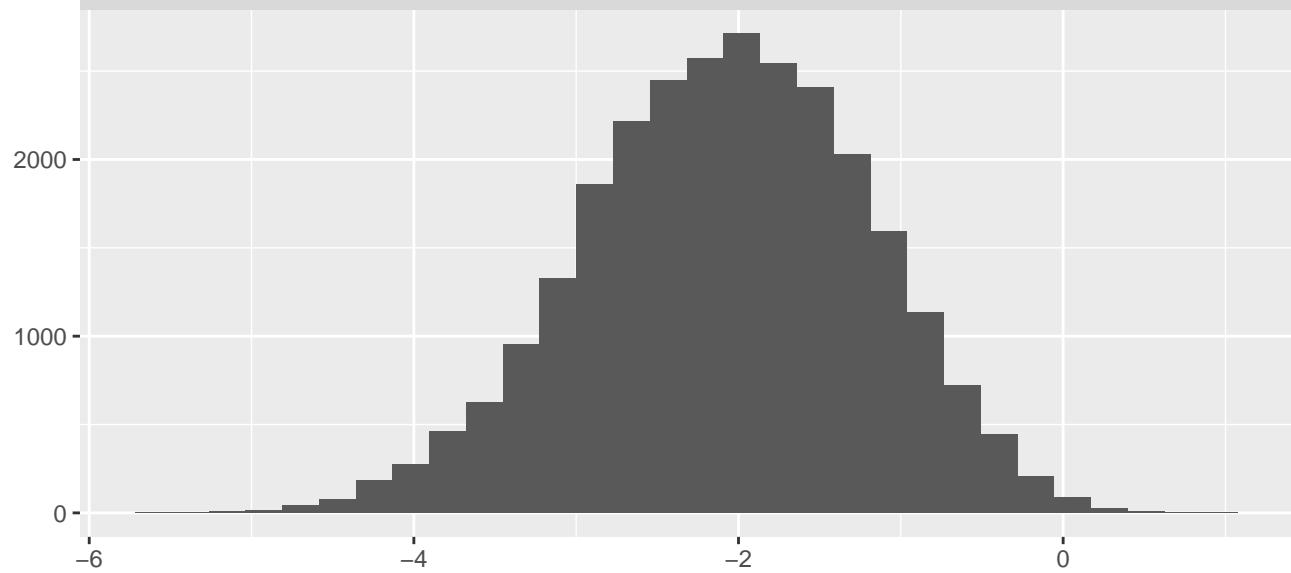
spline14



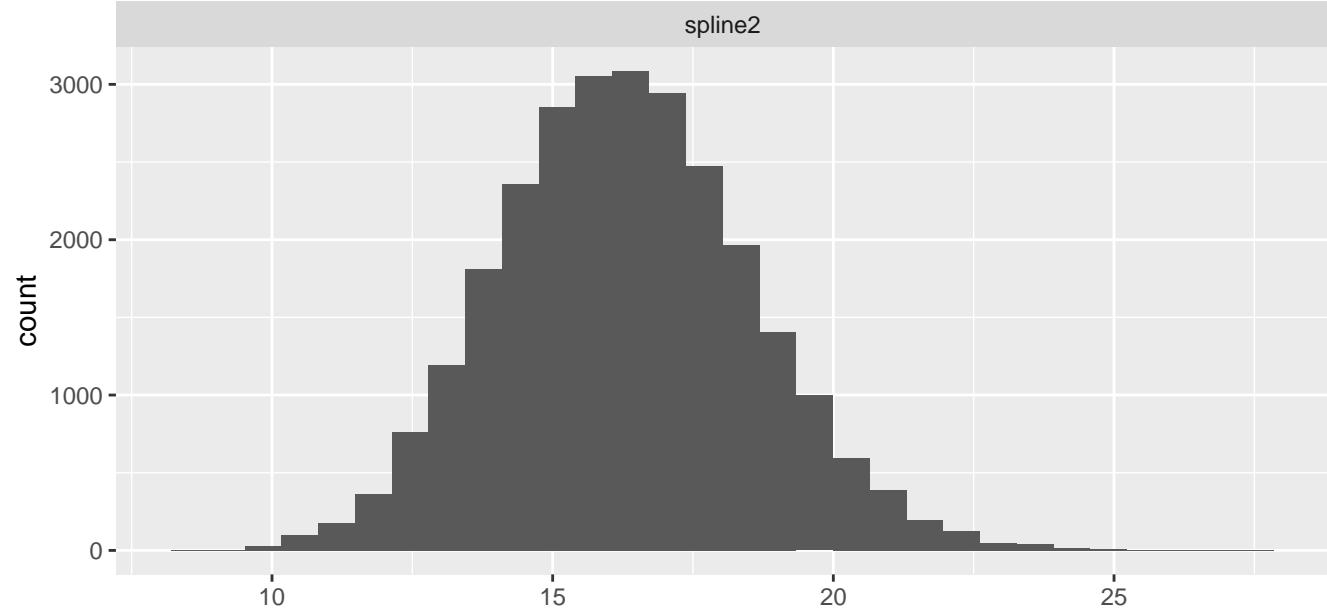
spline15



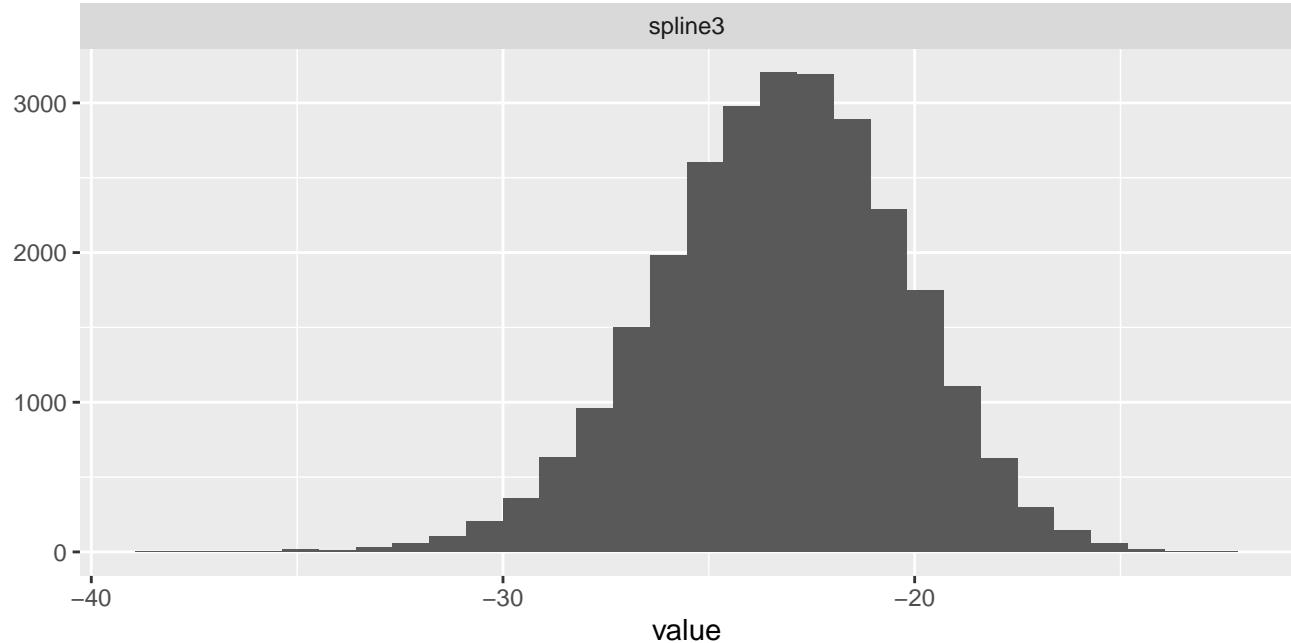
spline16



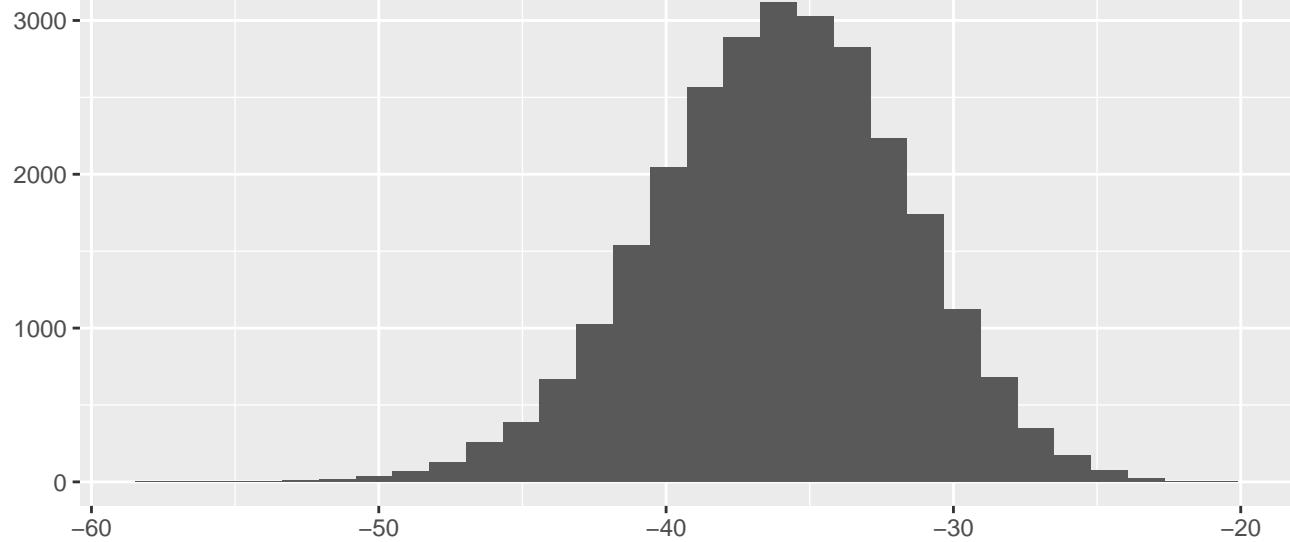
spline2



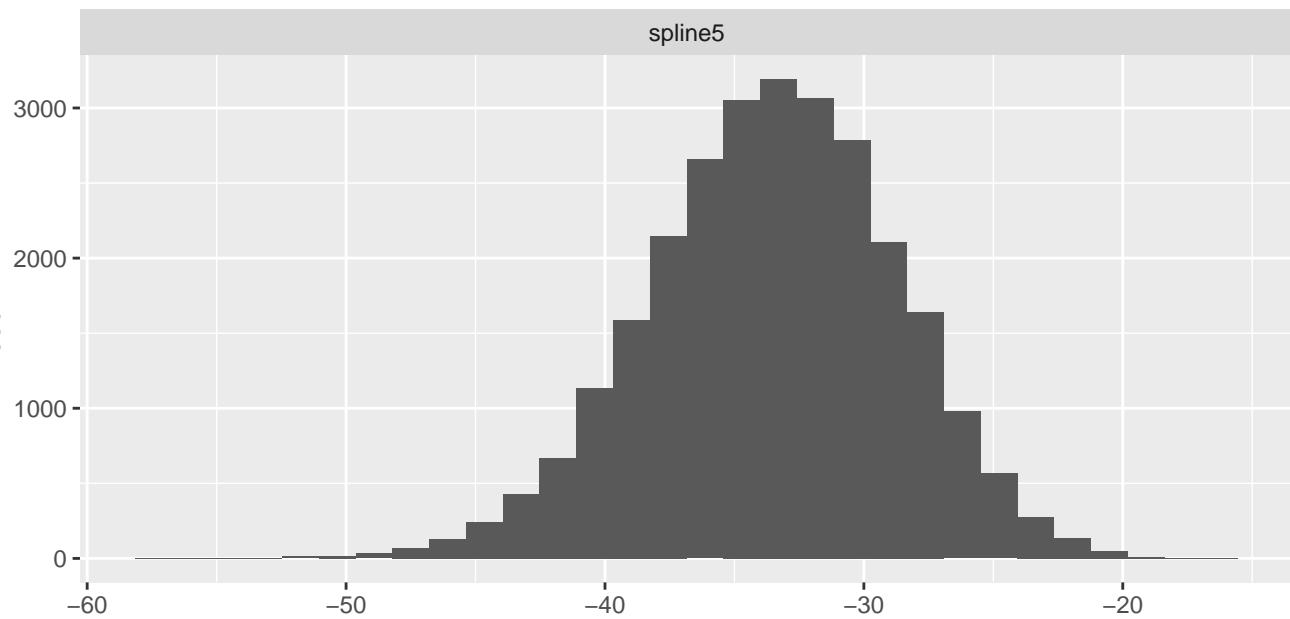
spline3



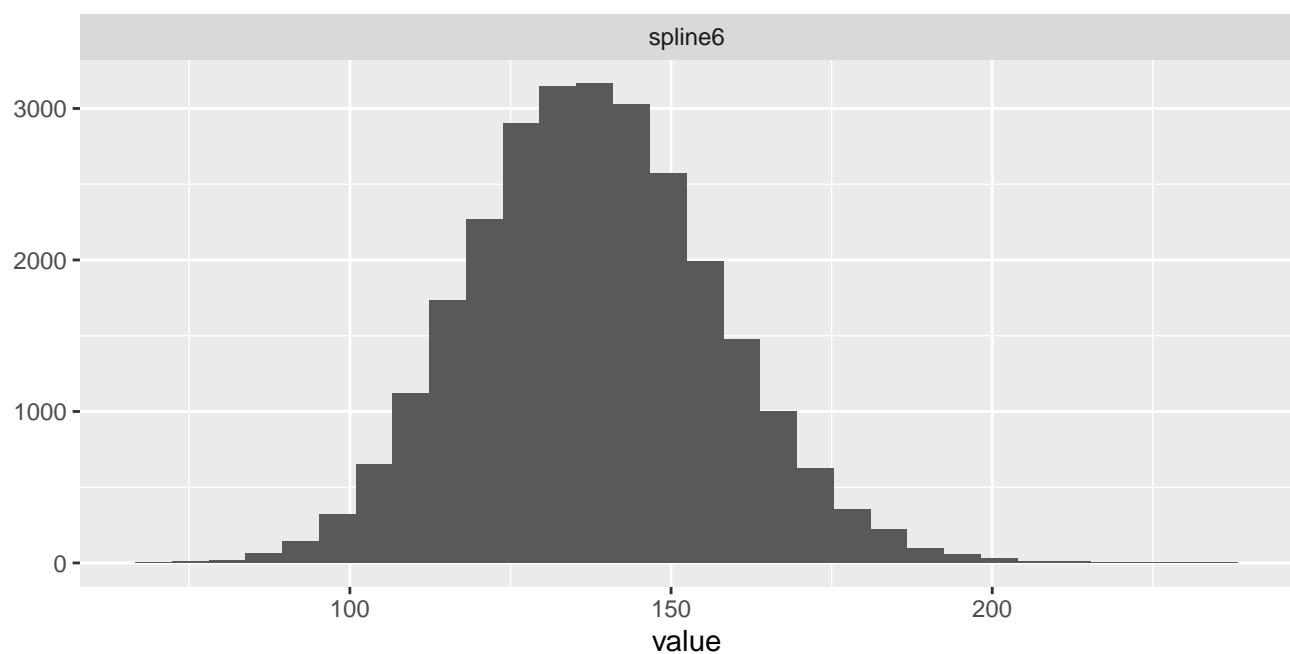
spline4



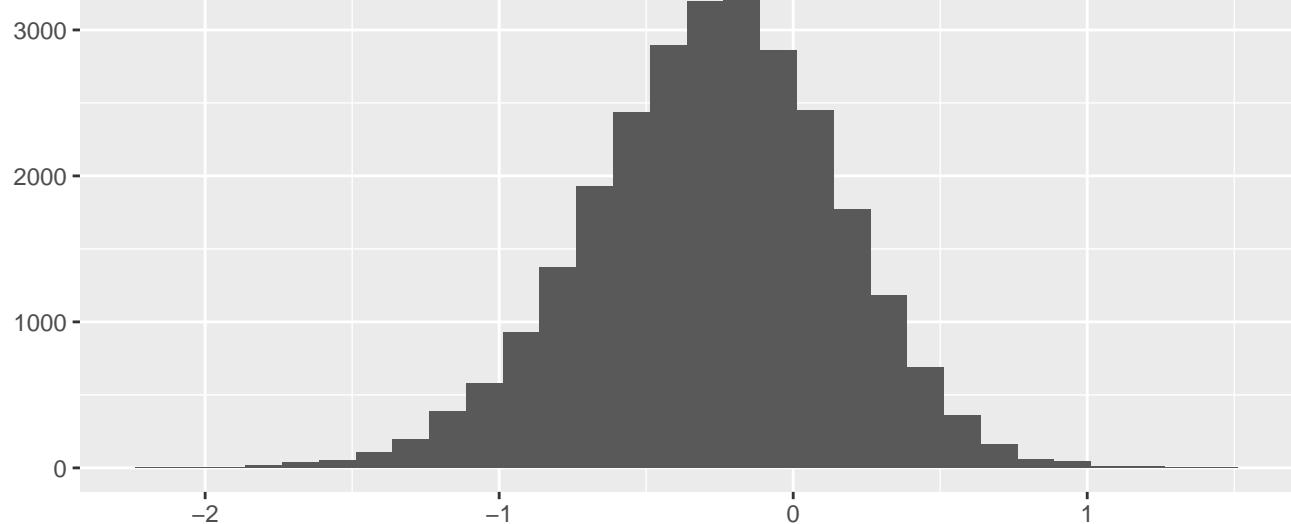
spline5



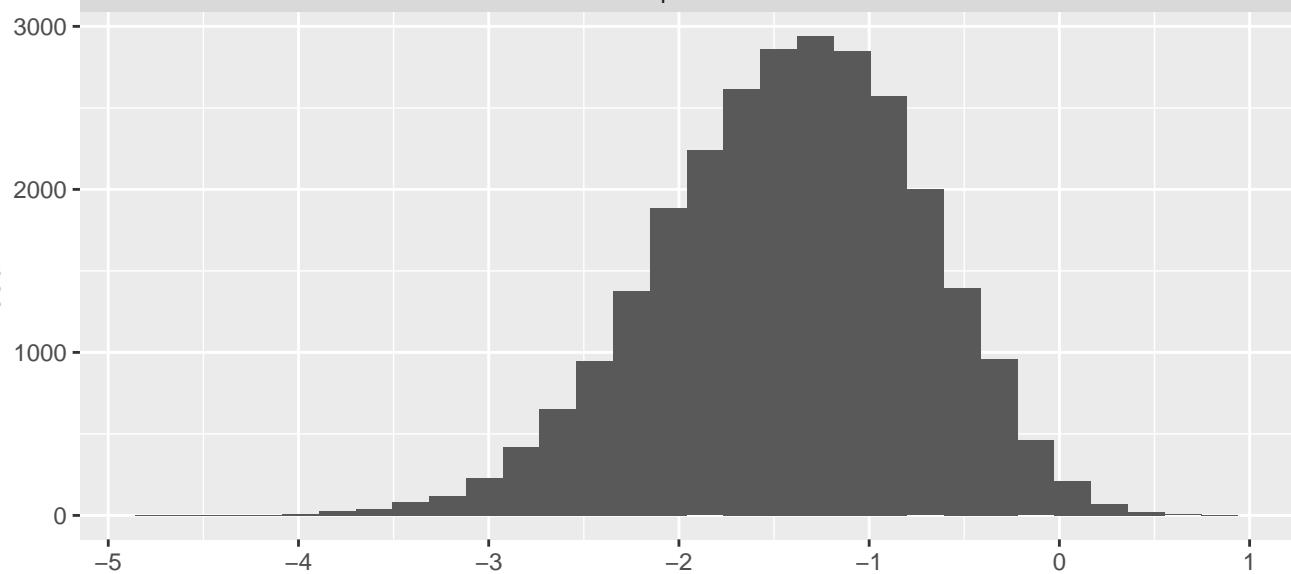
spline6



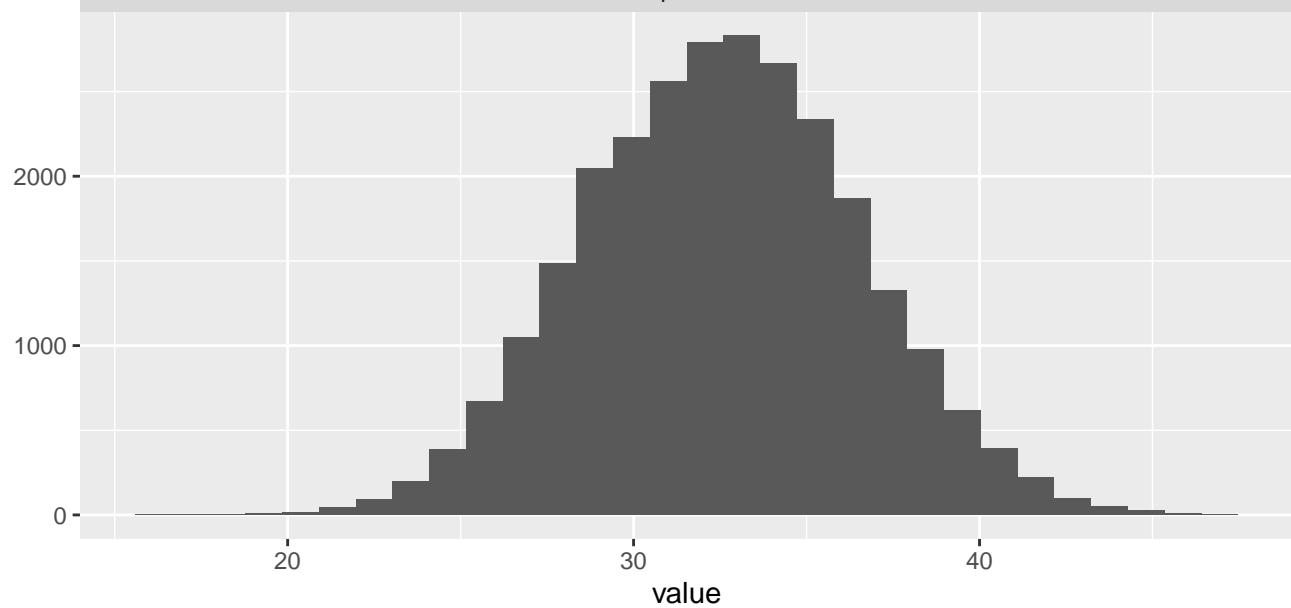
spline7



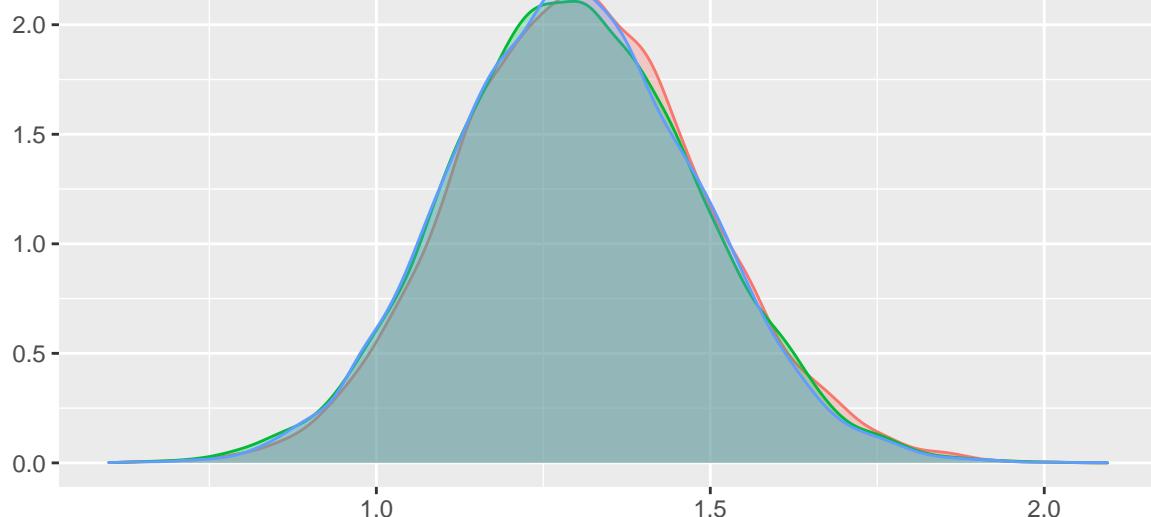
spline8



spline9



env.bio_15



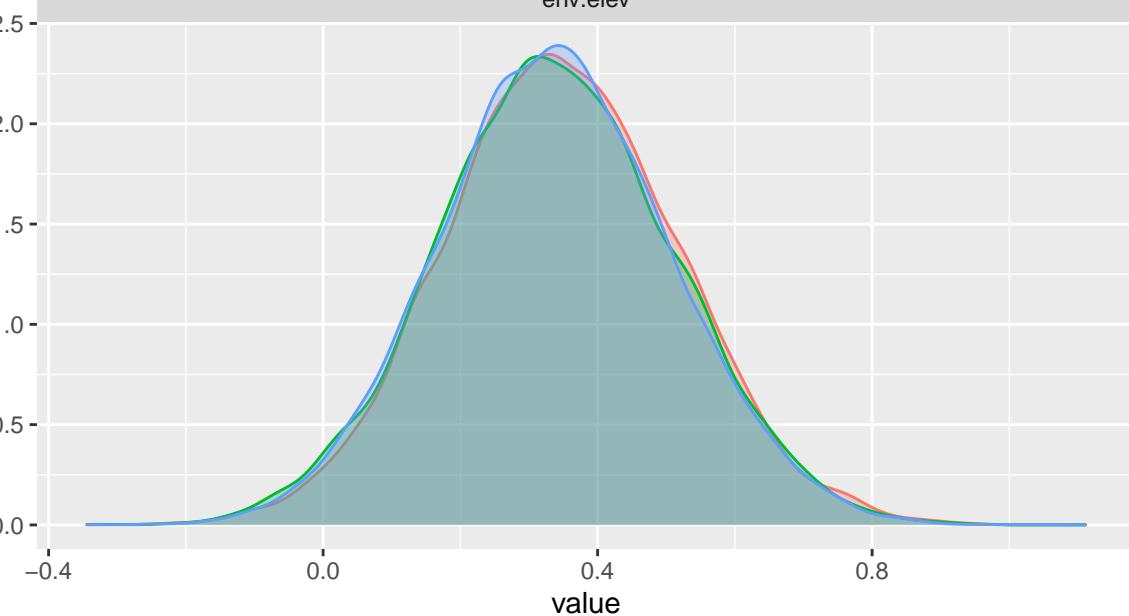
env.bio_7

density

Chain
1
2
3

-1.0 -0.8 -0.6 -0.4

env.elev



env.npp

4
2
0

0.8 1.0 1.2 1.4

Intercept

2.0
1.5
1.0
0.5
0.0

-24.0 -23.5 -23.0

Chain
1
2
3

spline1

0.10
0.05
0.00

20 30

value

spline10

0.10
0.05
0.00

20 30

spline11

0.100
0.075
0.050
0.025
0.000

-50 -40 -30 -20

Chain
1
2
3

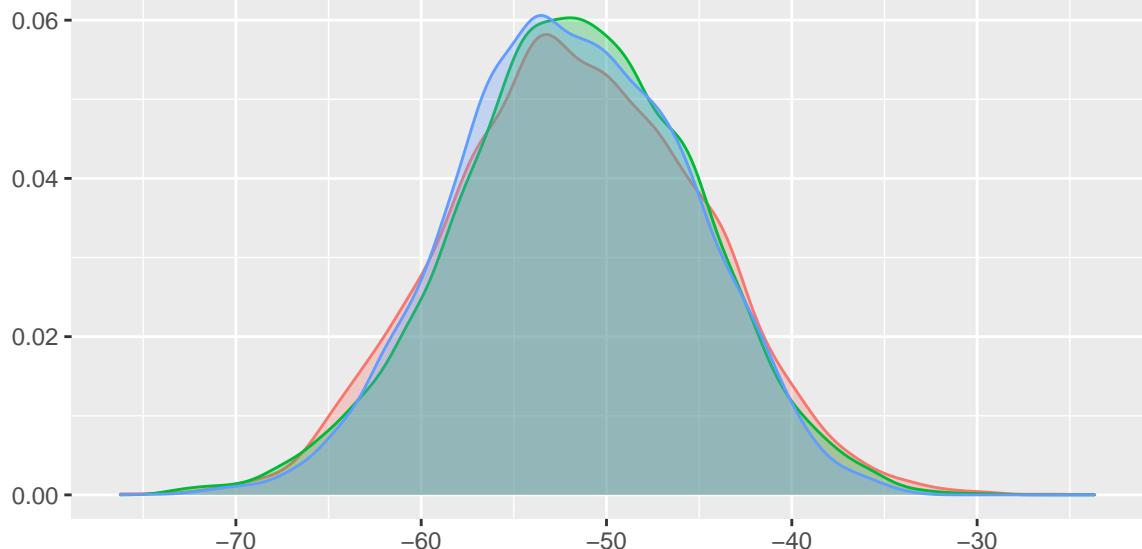
spline12

0.06
0.04
0.02
0.00

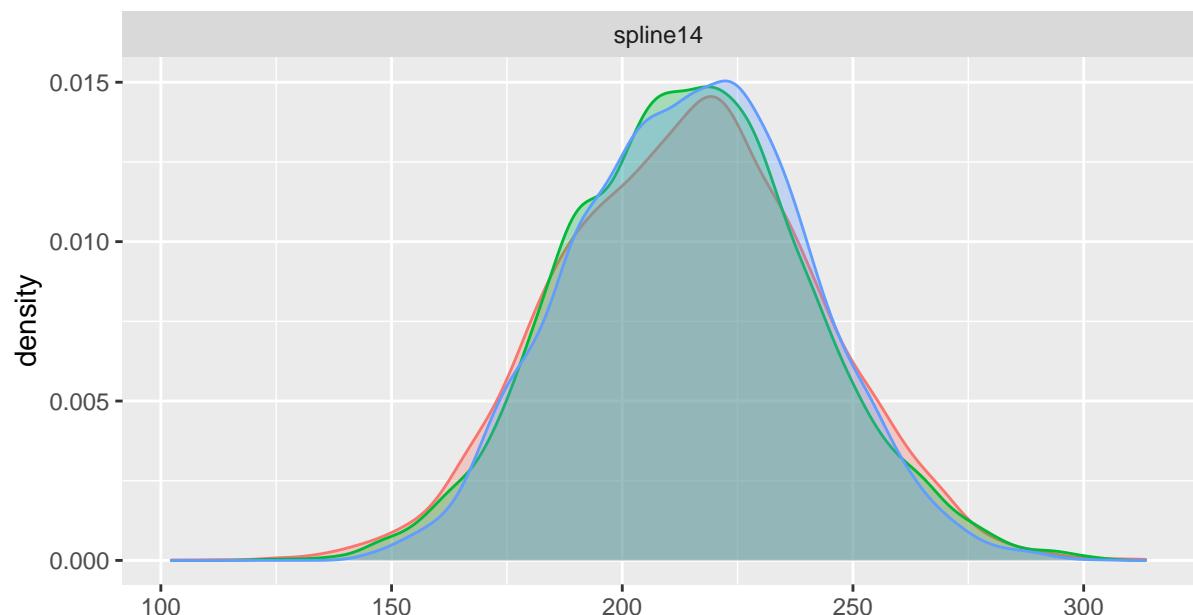
-70 -60 -50 -40 -30

value

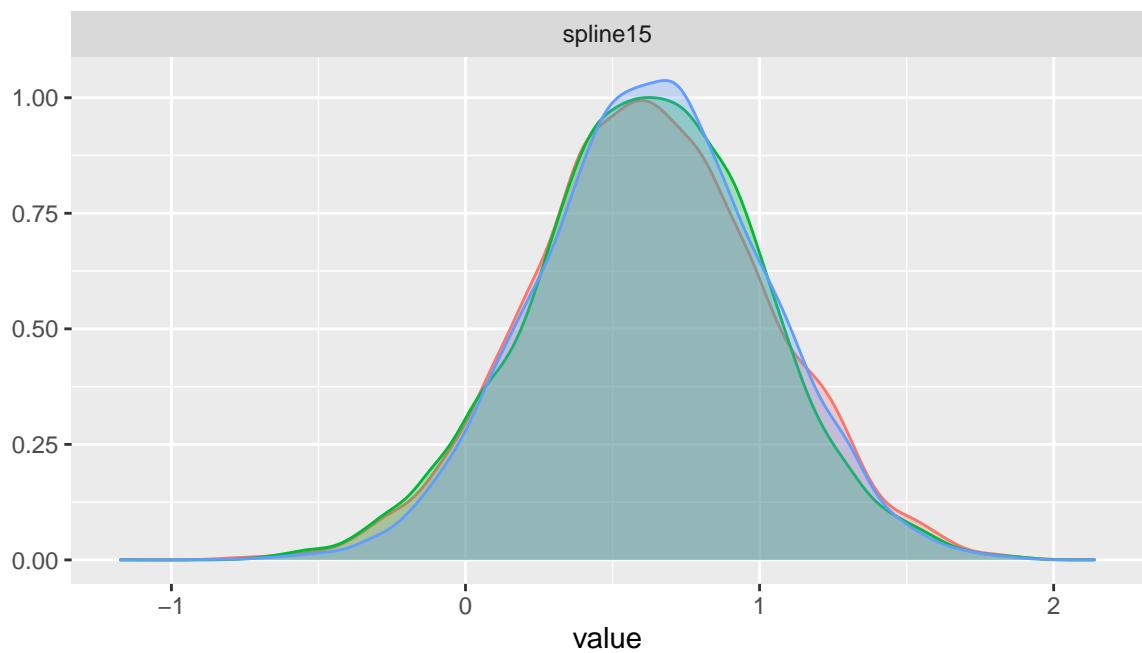
spline13



spline14

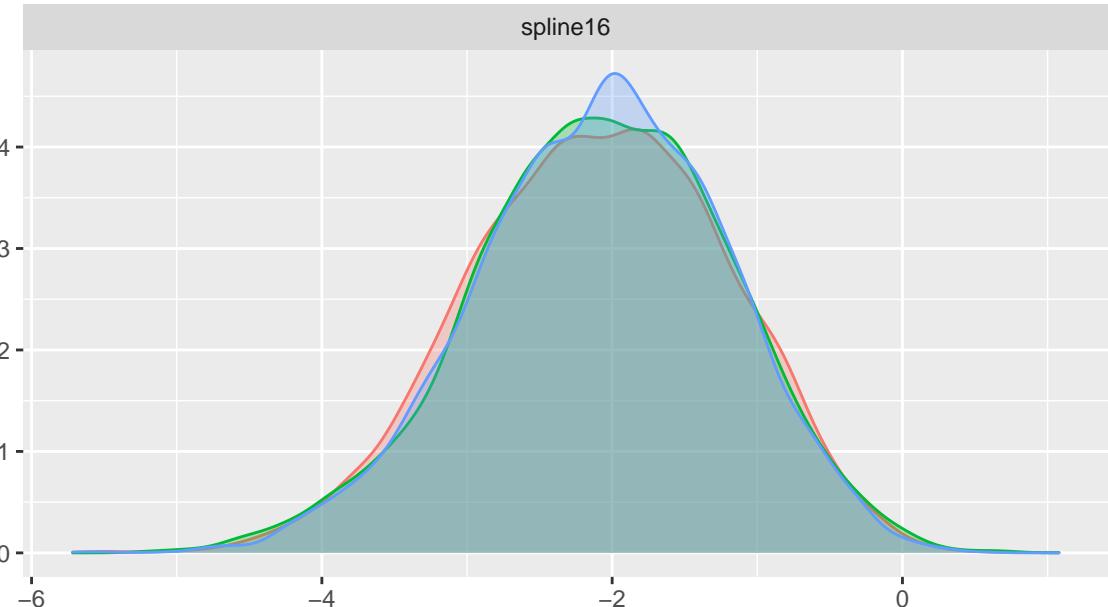


spline15

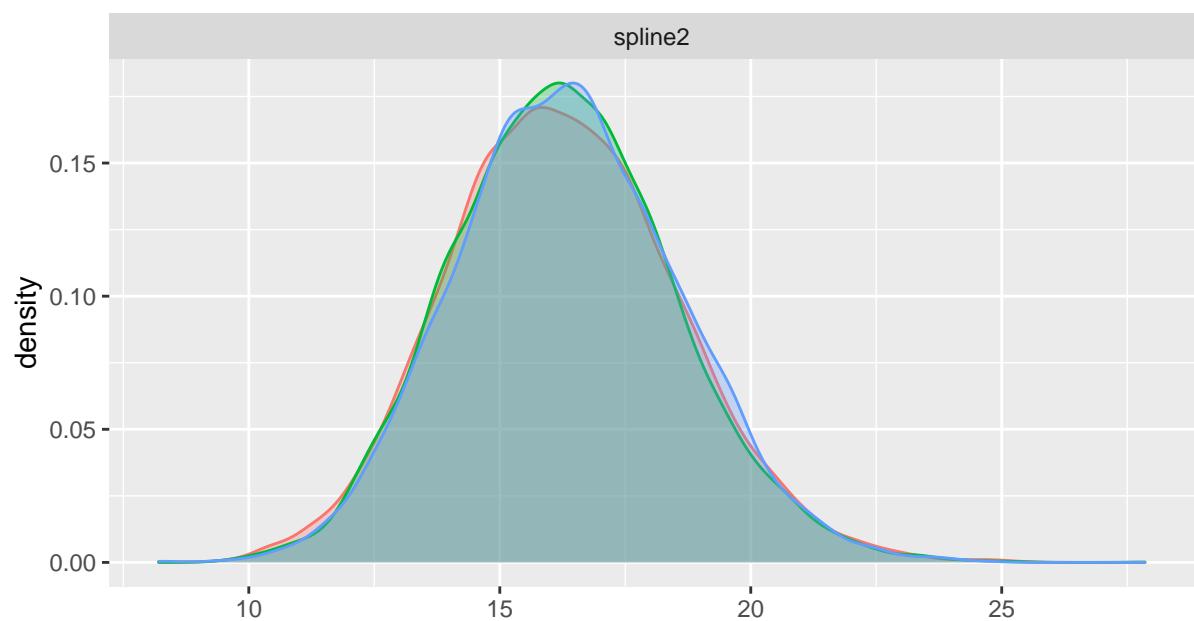


Chain
1
2
3

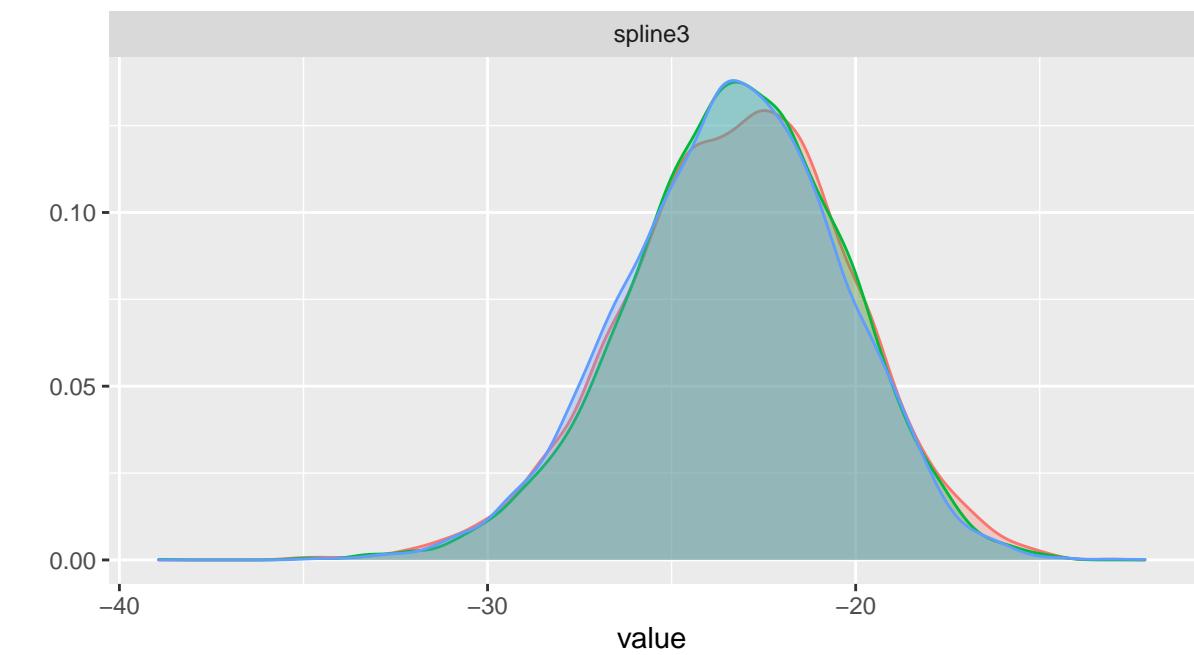
spline16



spline2

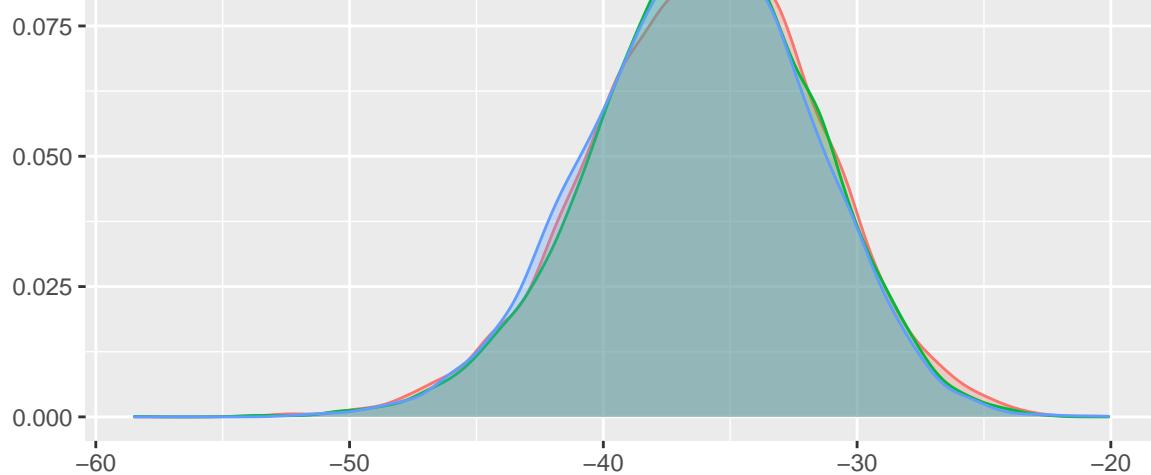


spline3

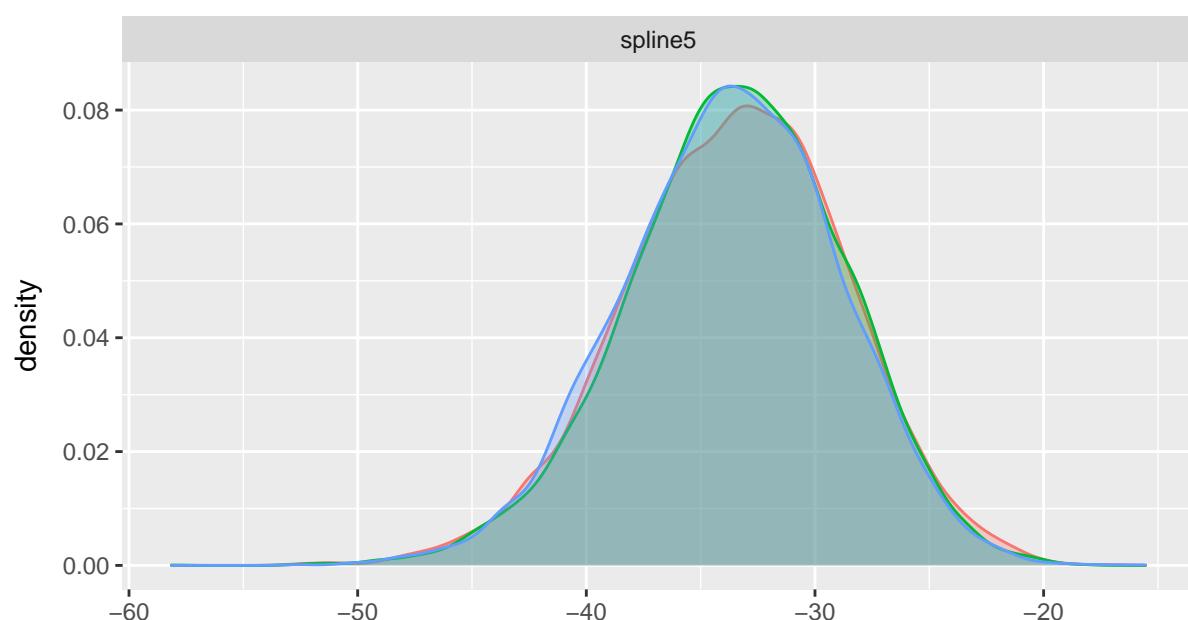


Chain
1
2
3

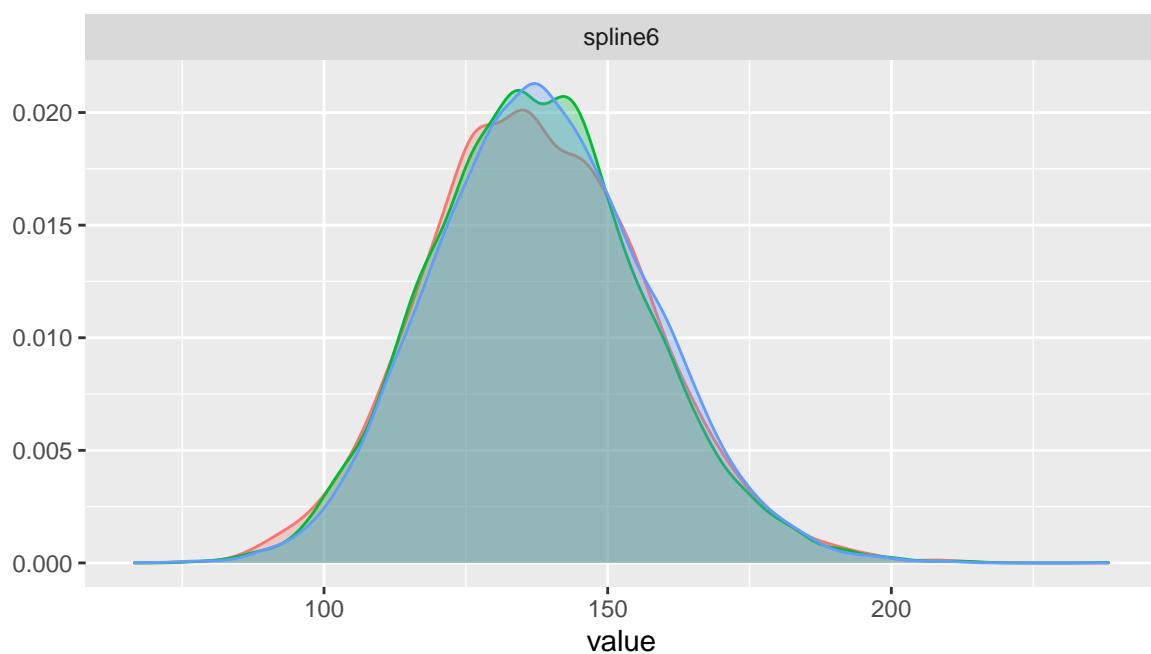
spline4



spline5

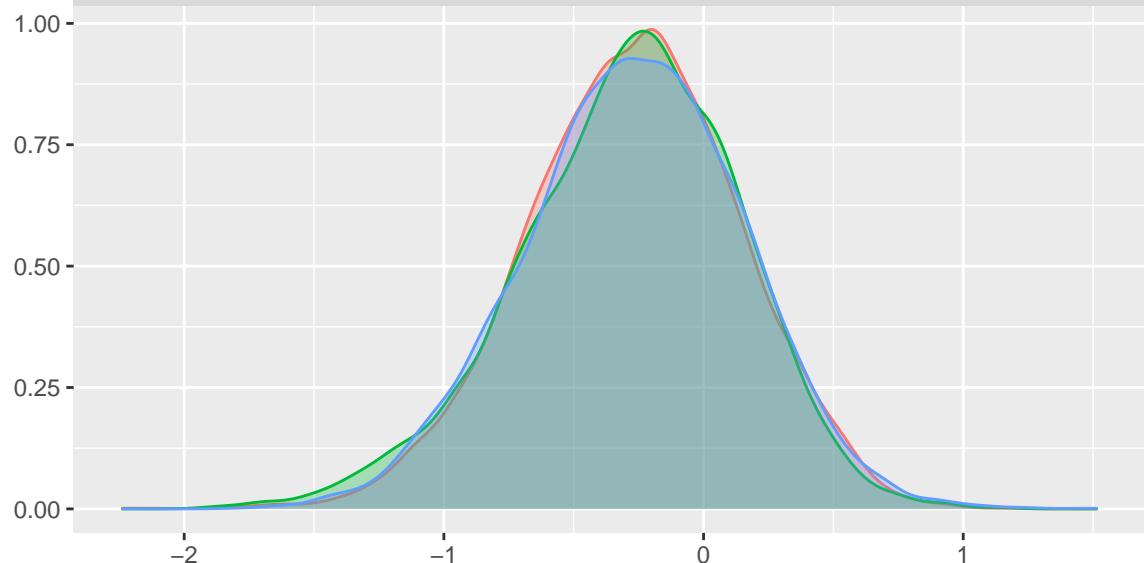


spline6

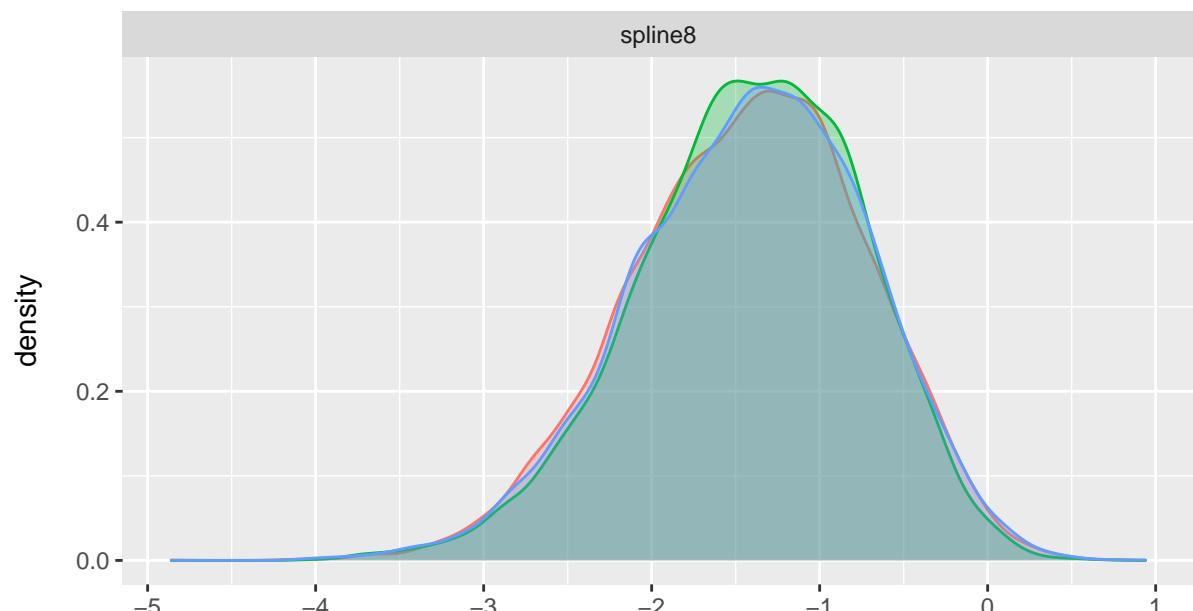


Chain
1
2
3

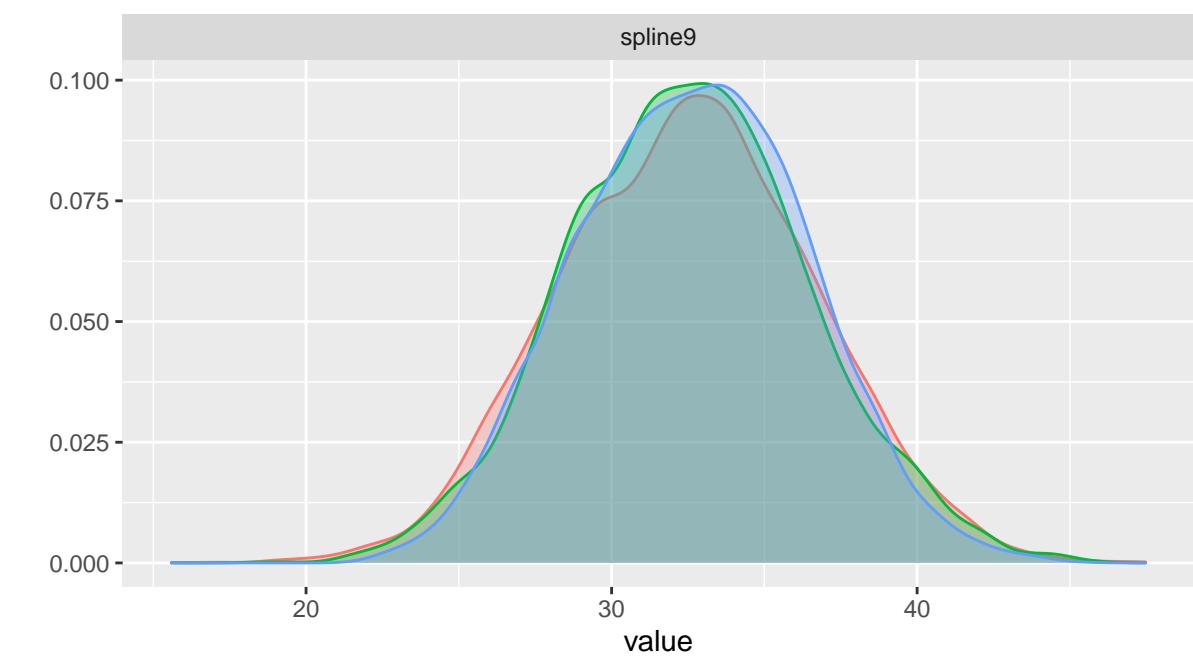
spline7



spline8

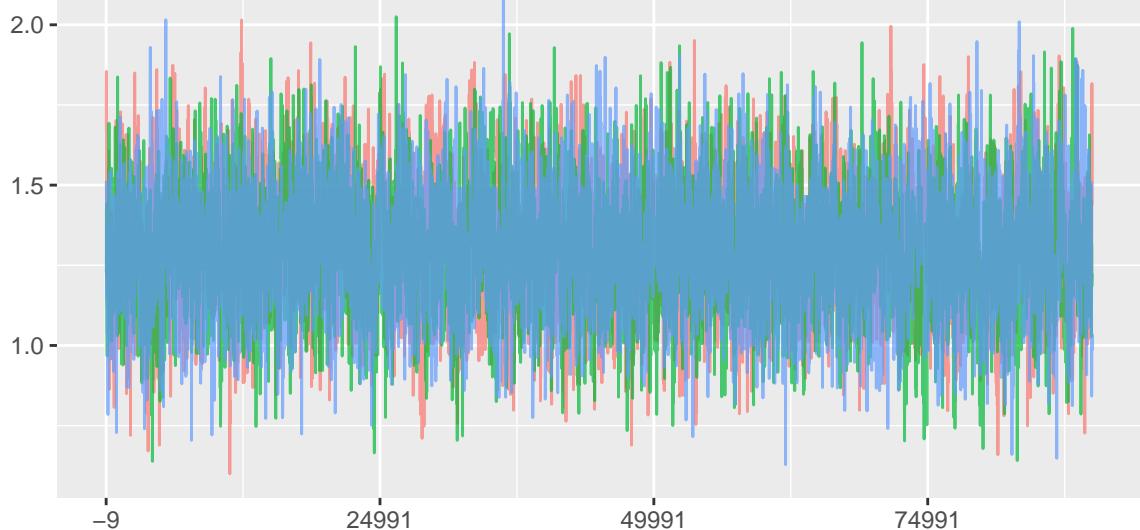


spline9

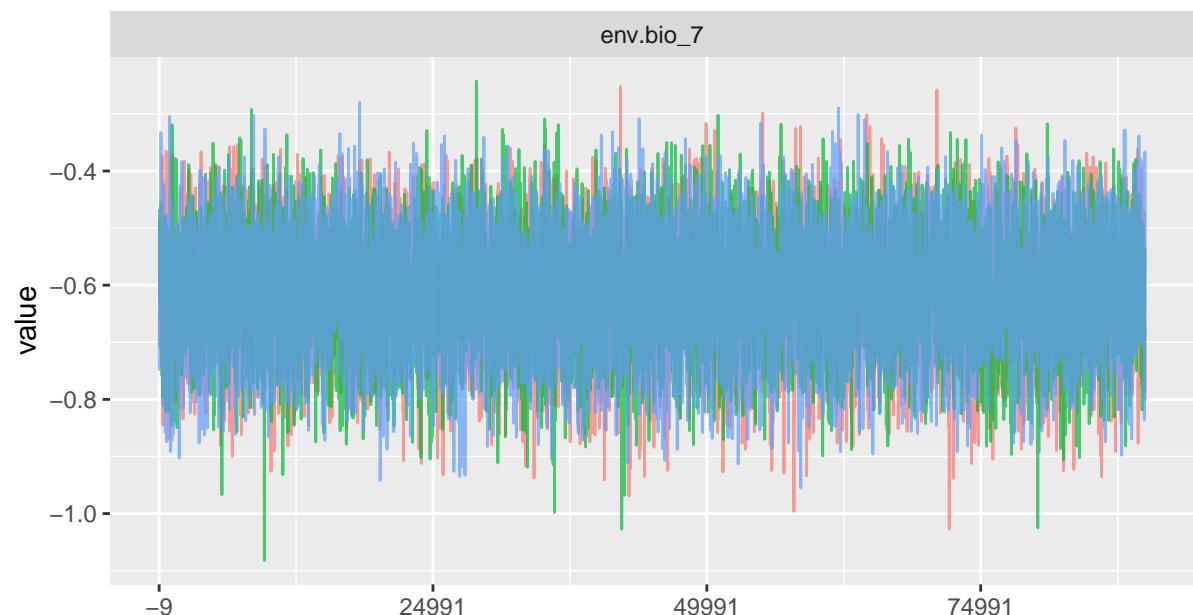


Chain
1
2
3

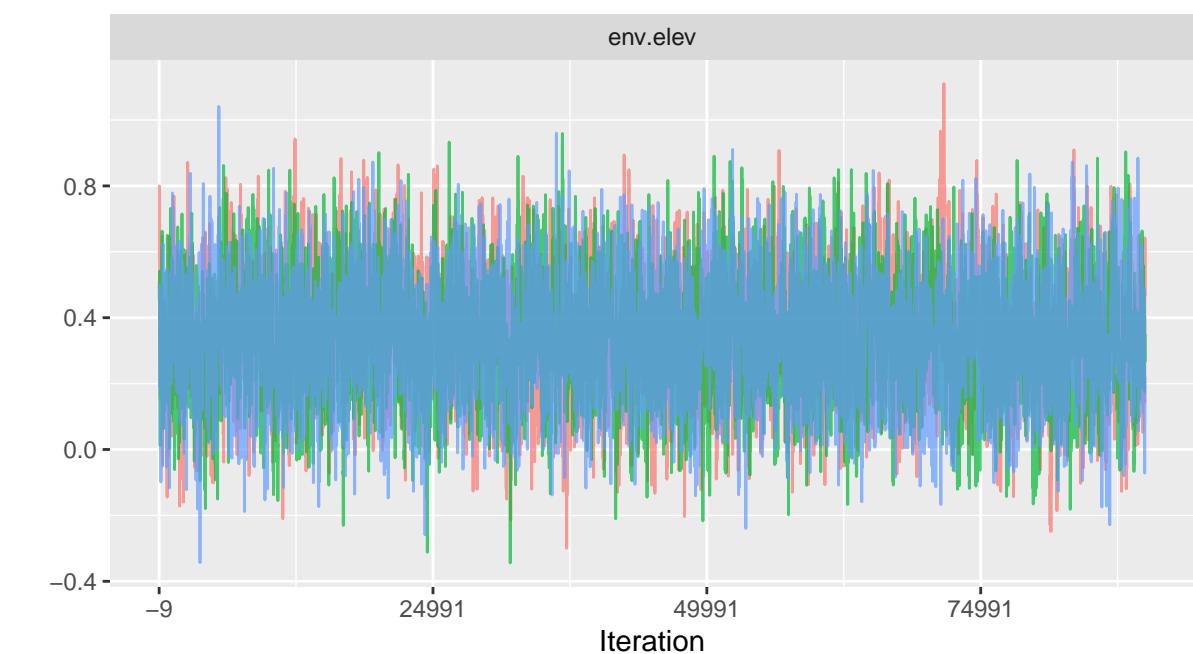
env.bio_15



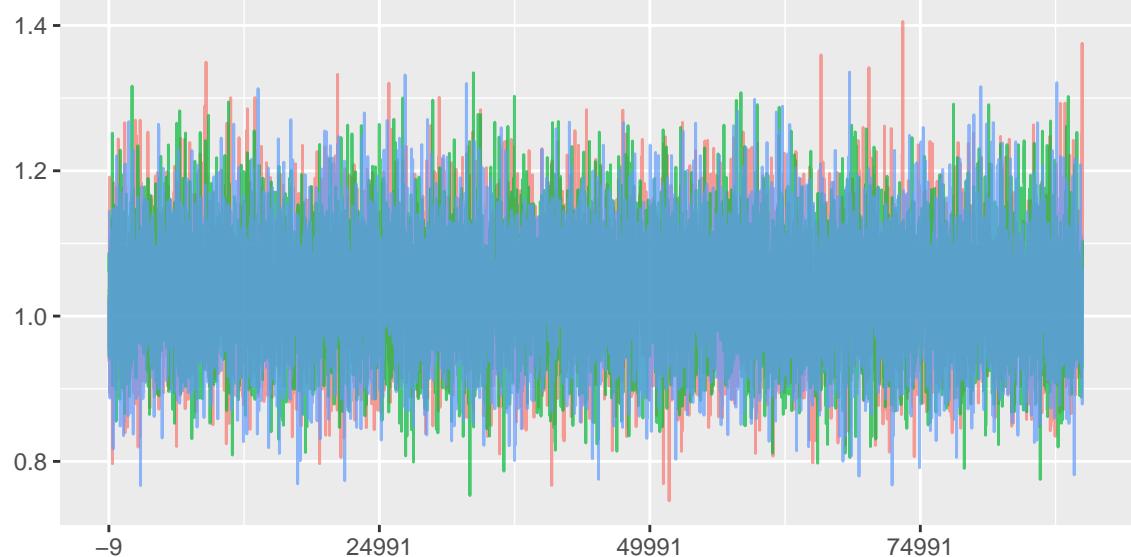
env.bio_7



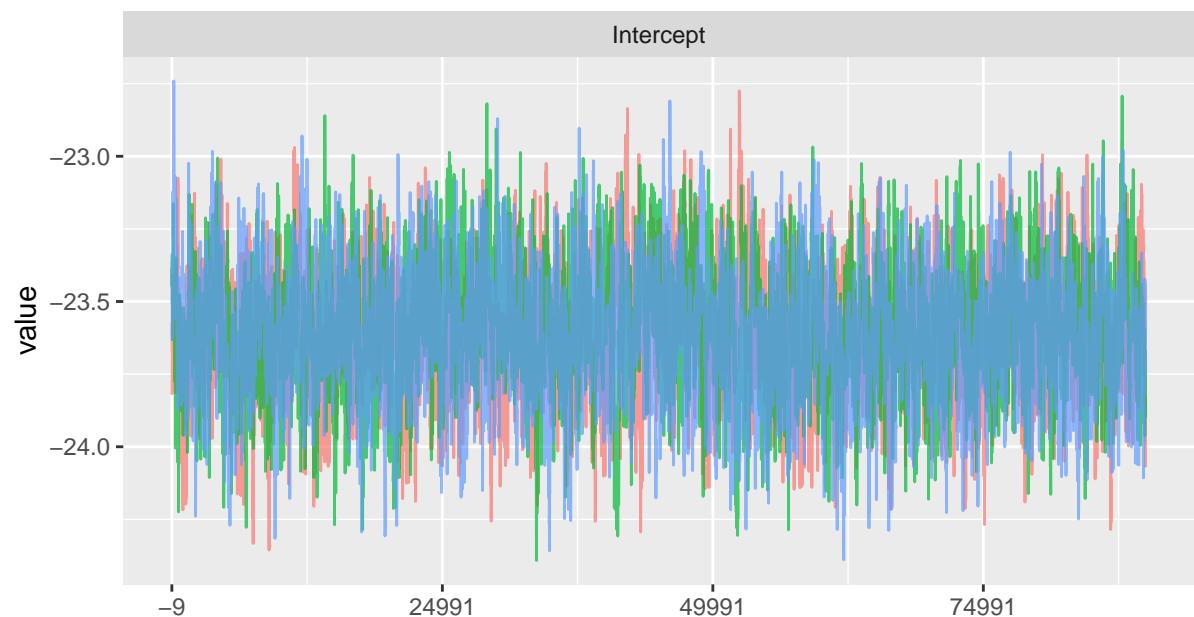
env.elev



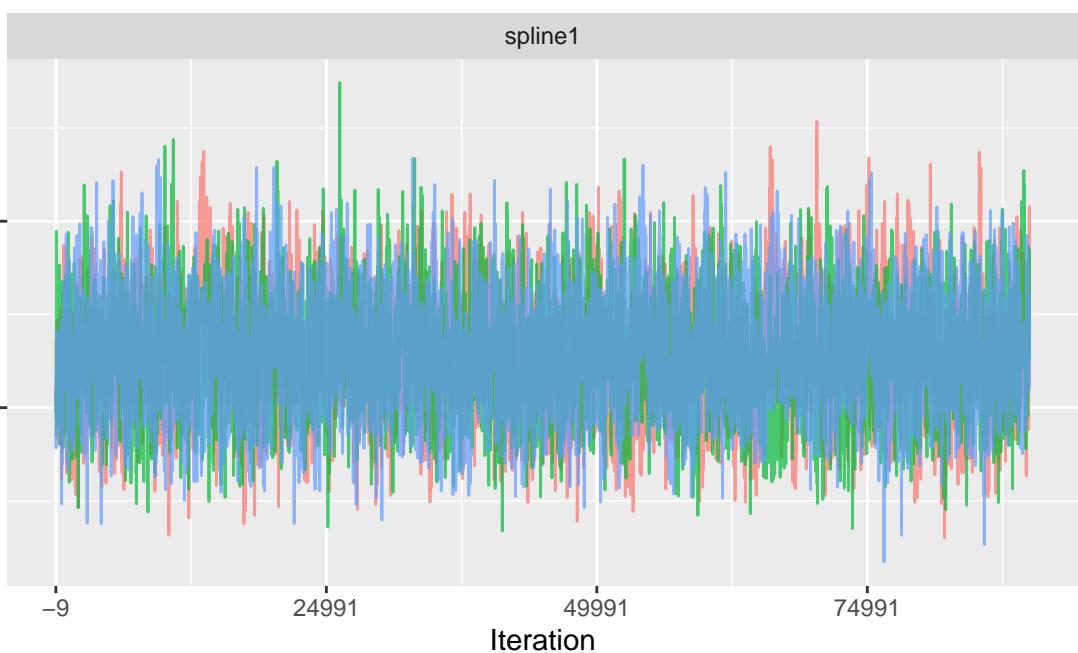
env.npp



Intercept



spline1

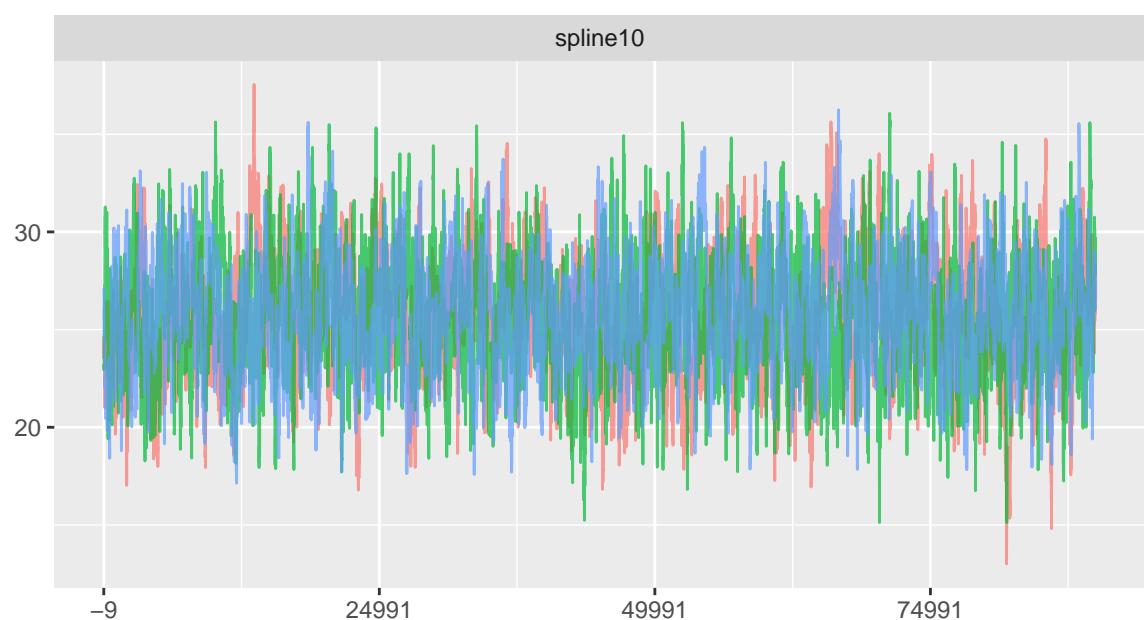


Iteration

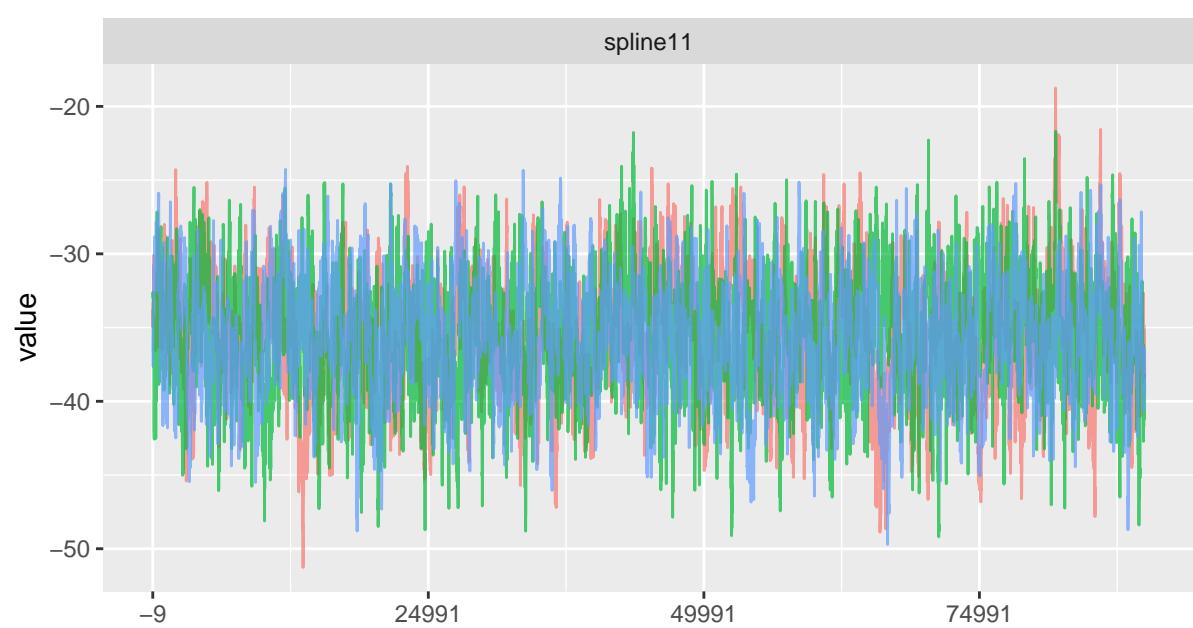
Chain

- 1
- 2
- 3

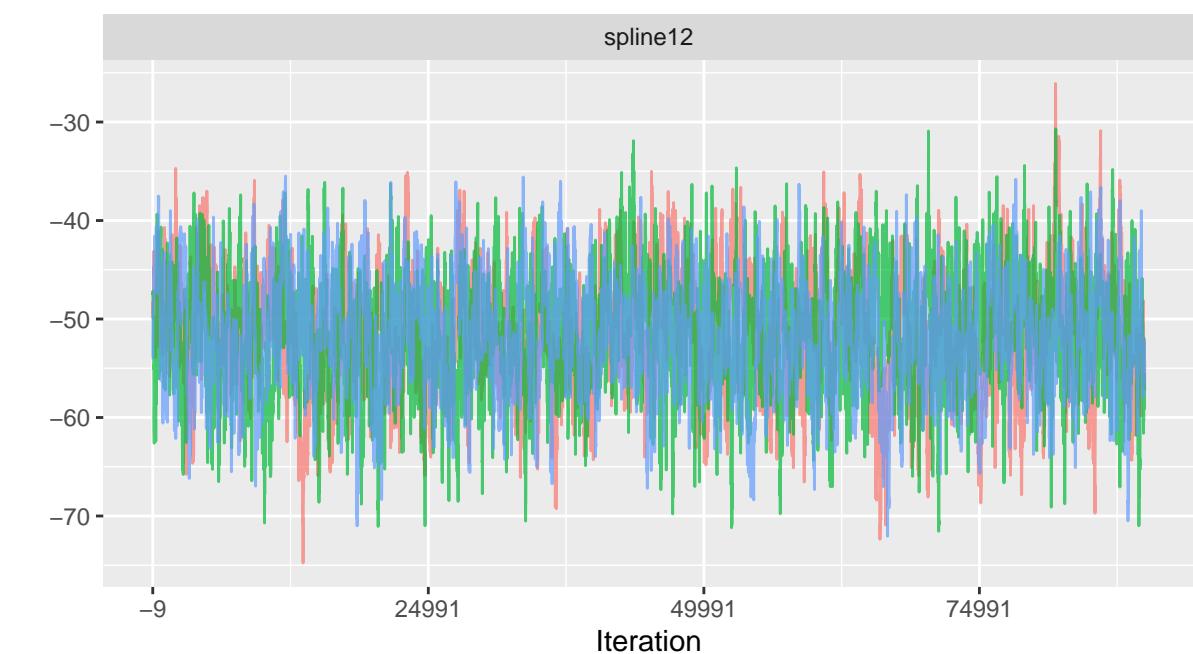
spline10



spline11



spline12

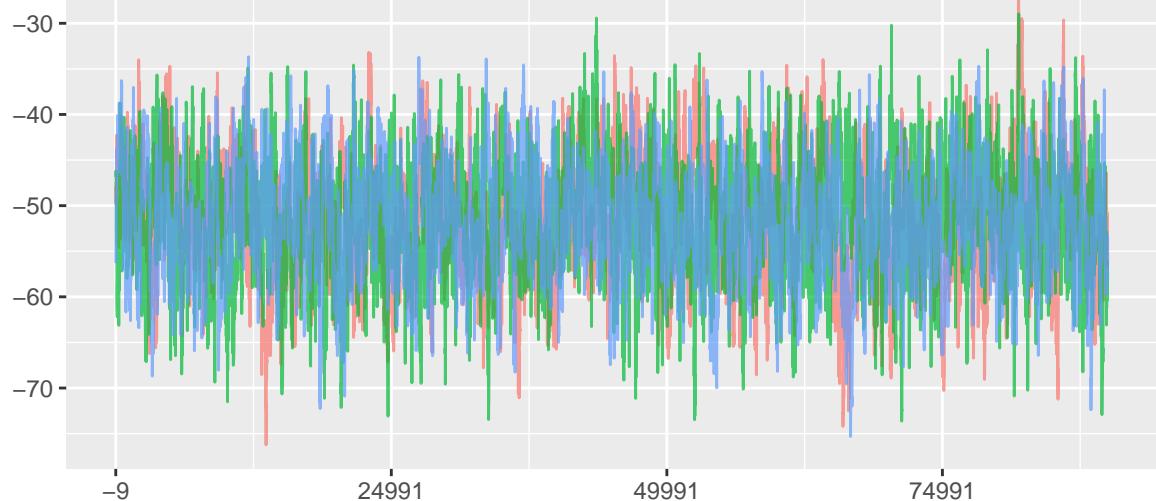


Chain

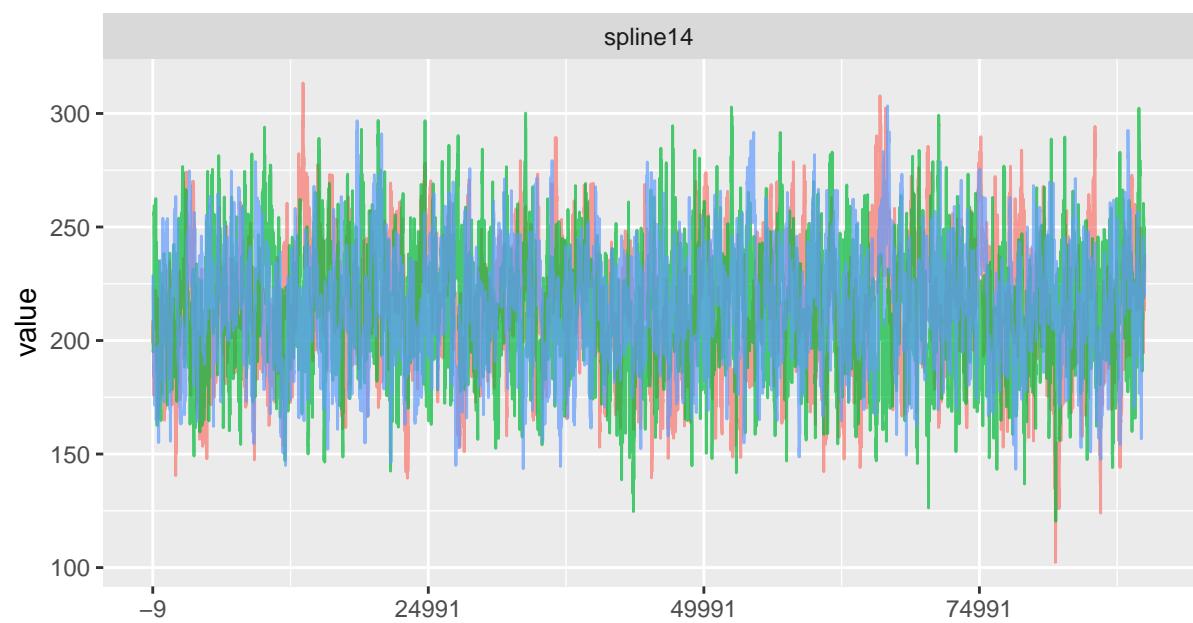
- 1
- 2
- 3

Iteration

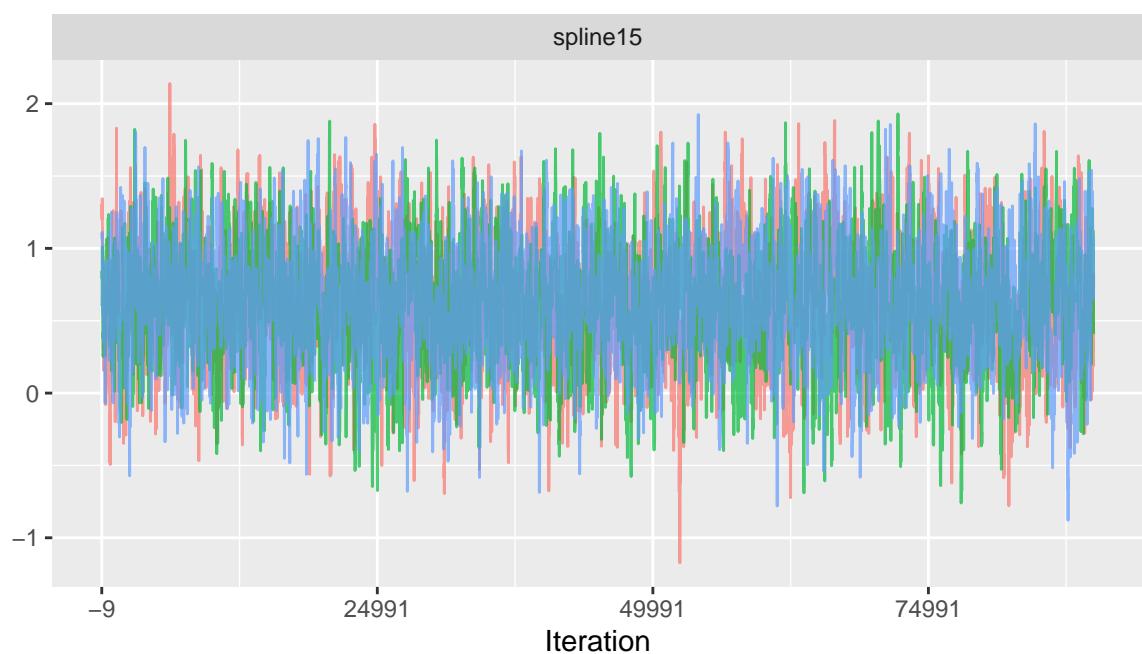
spline13



spline14



spline15

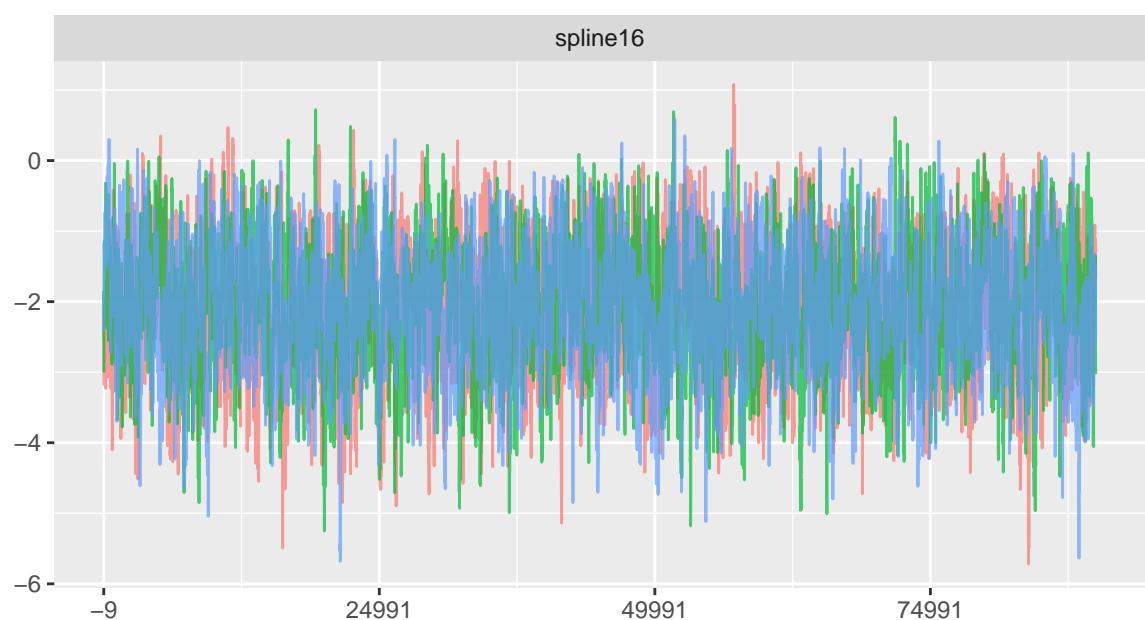


Iteration

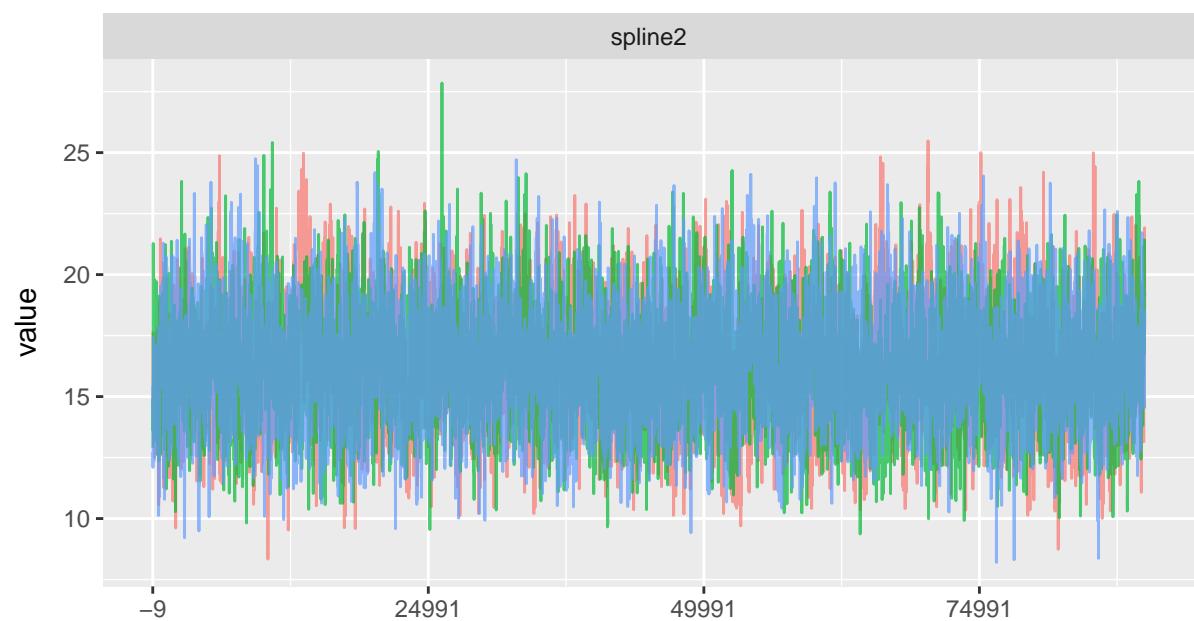
Chain

- 1
- 2
- 3

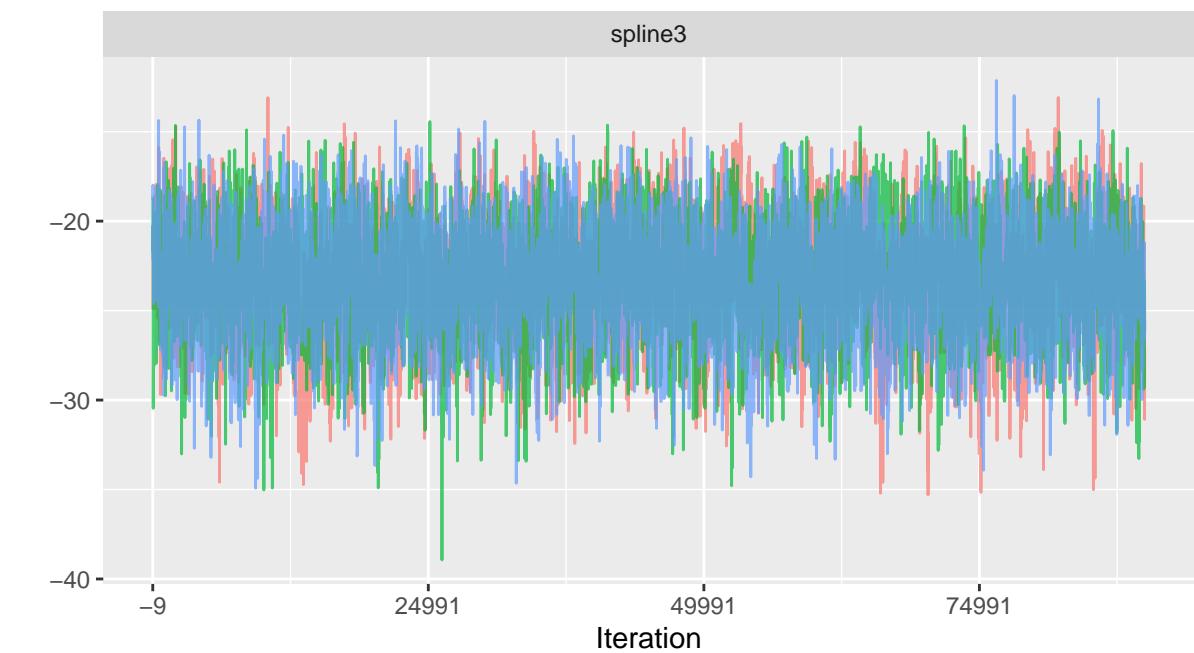
spline16



spline2



spline3

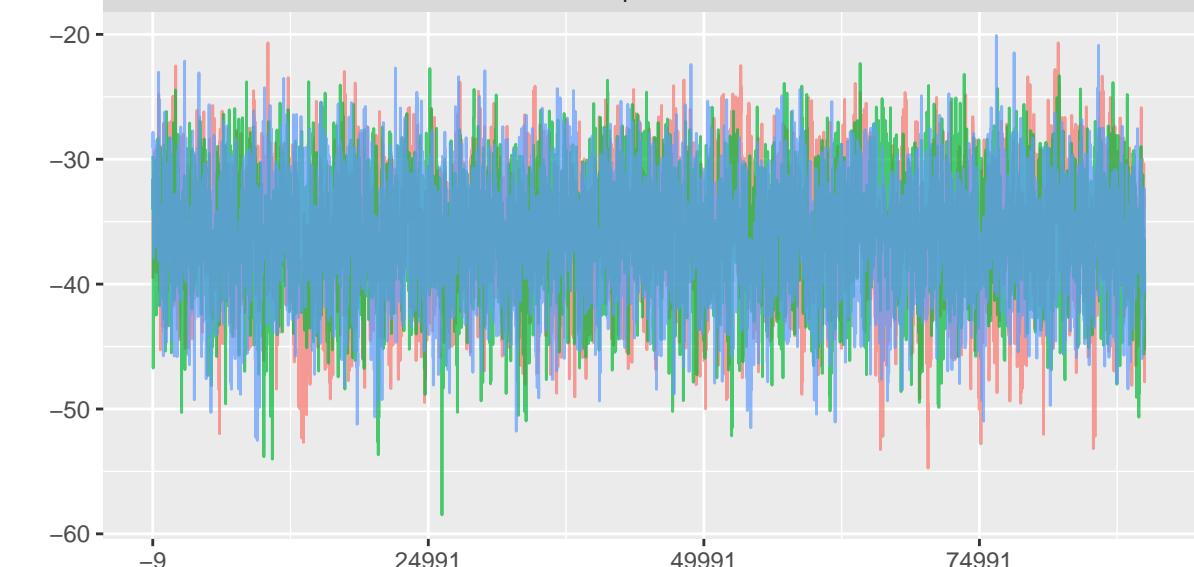


Chain

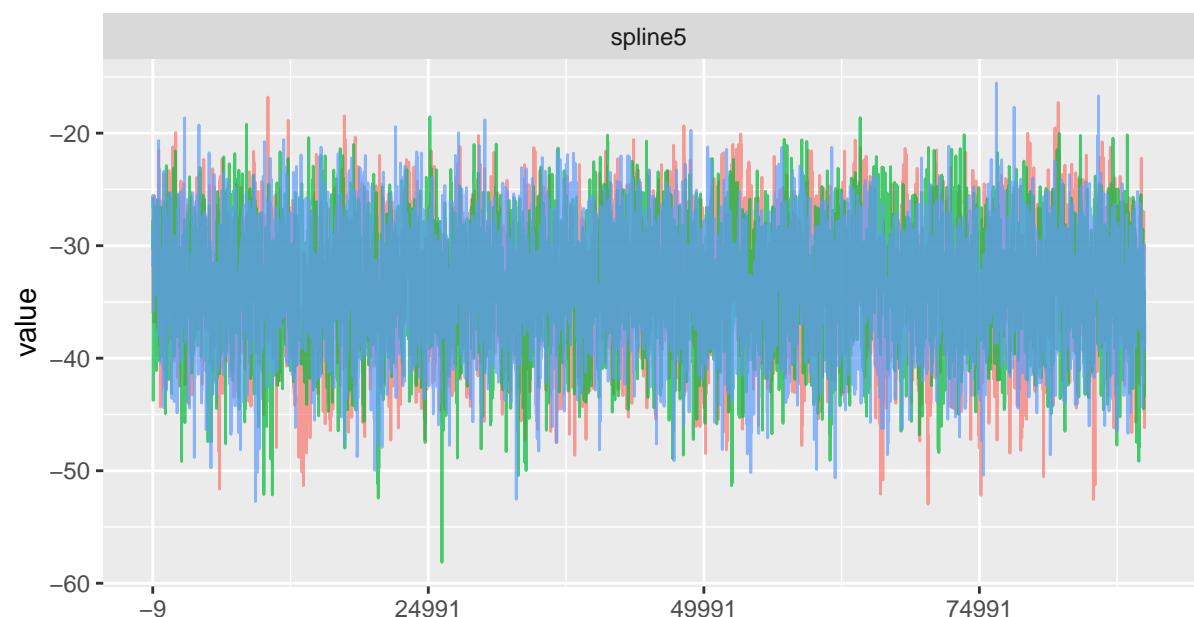
- 1
- 2
- 3

Iteration

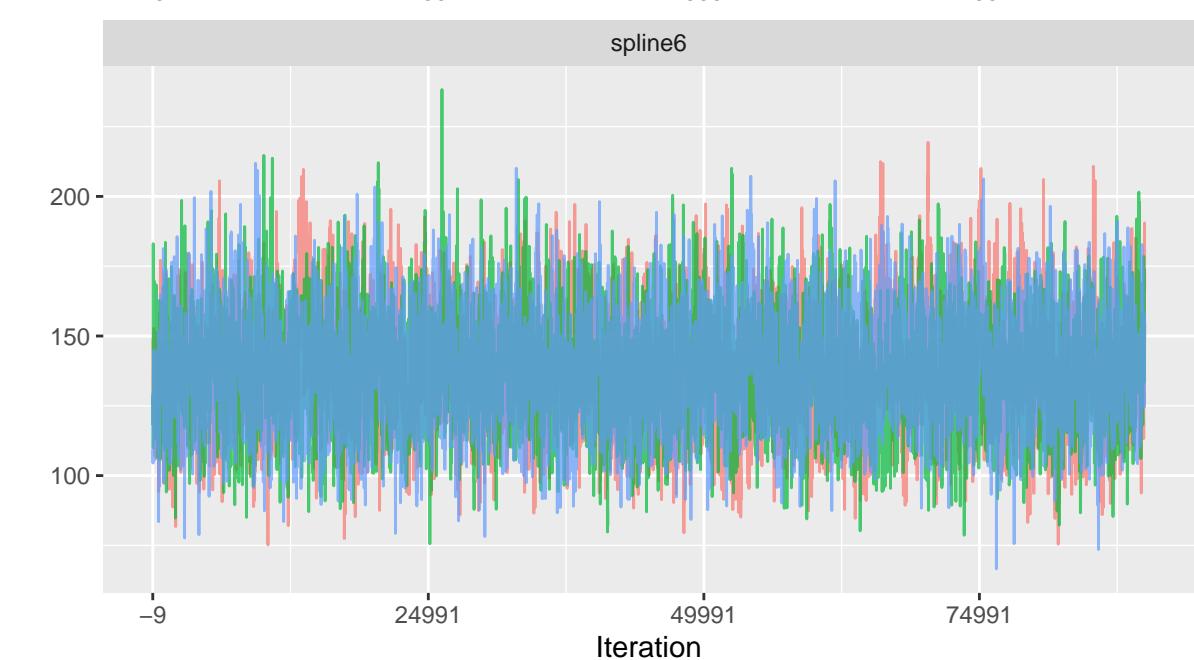
spline4



spline5



spline6

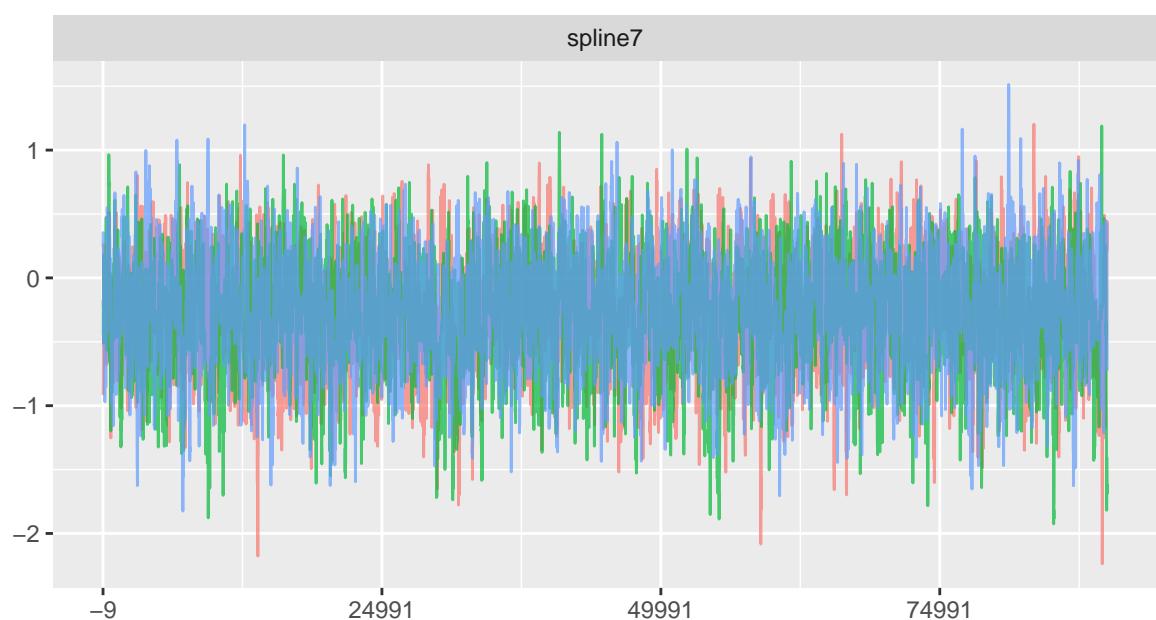


Chain

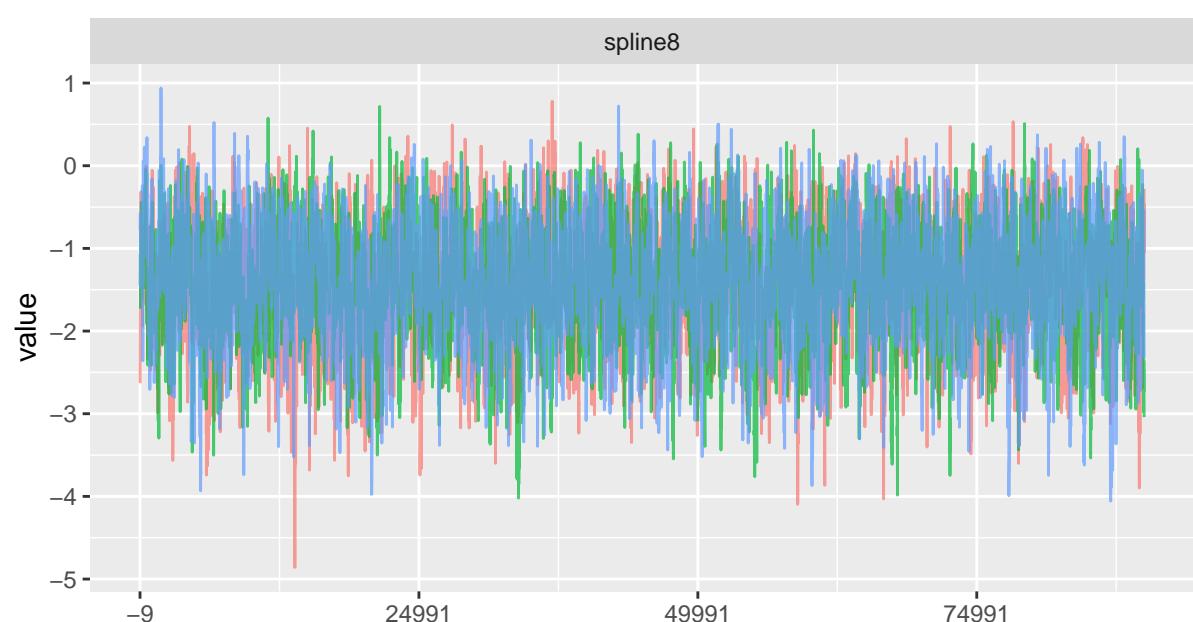
- 1
- 2
- 3

Iteration

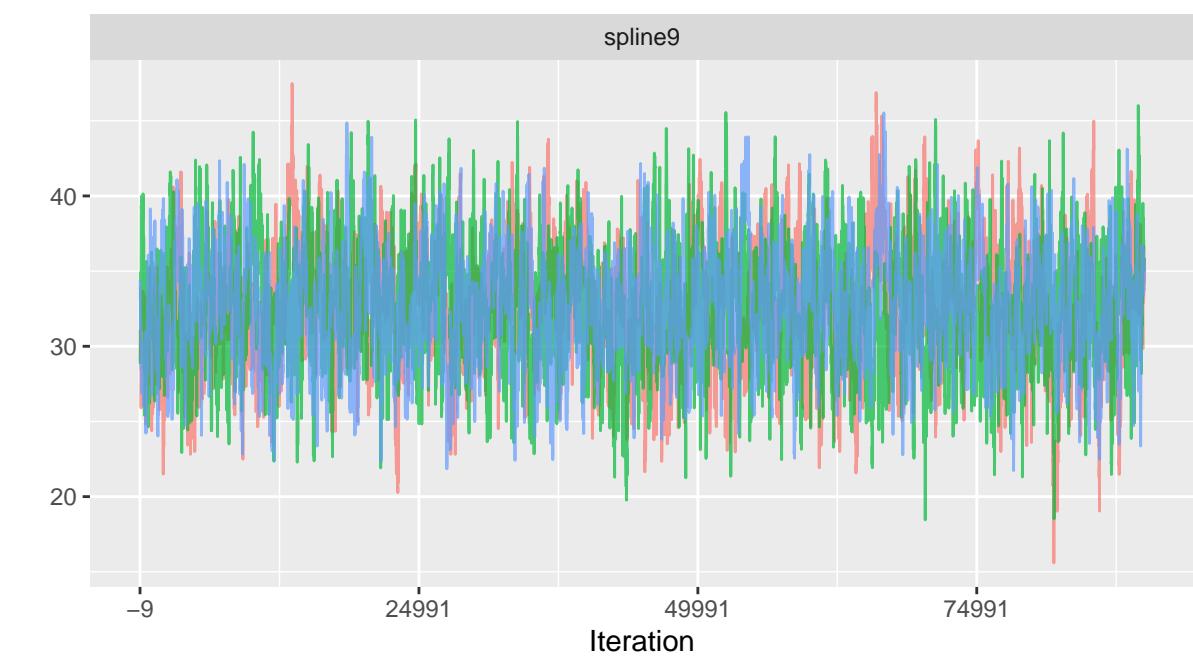
spline7



spline8



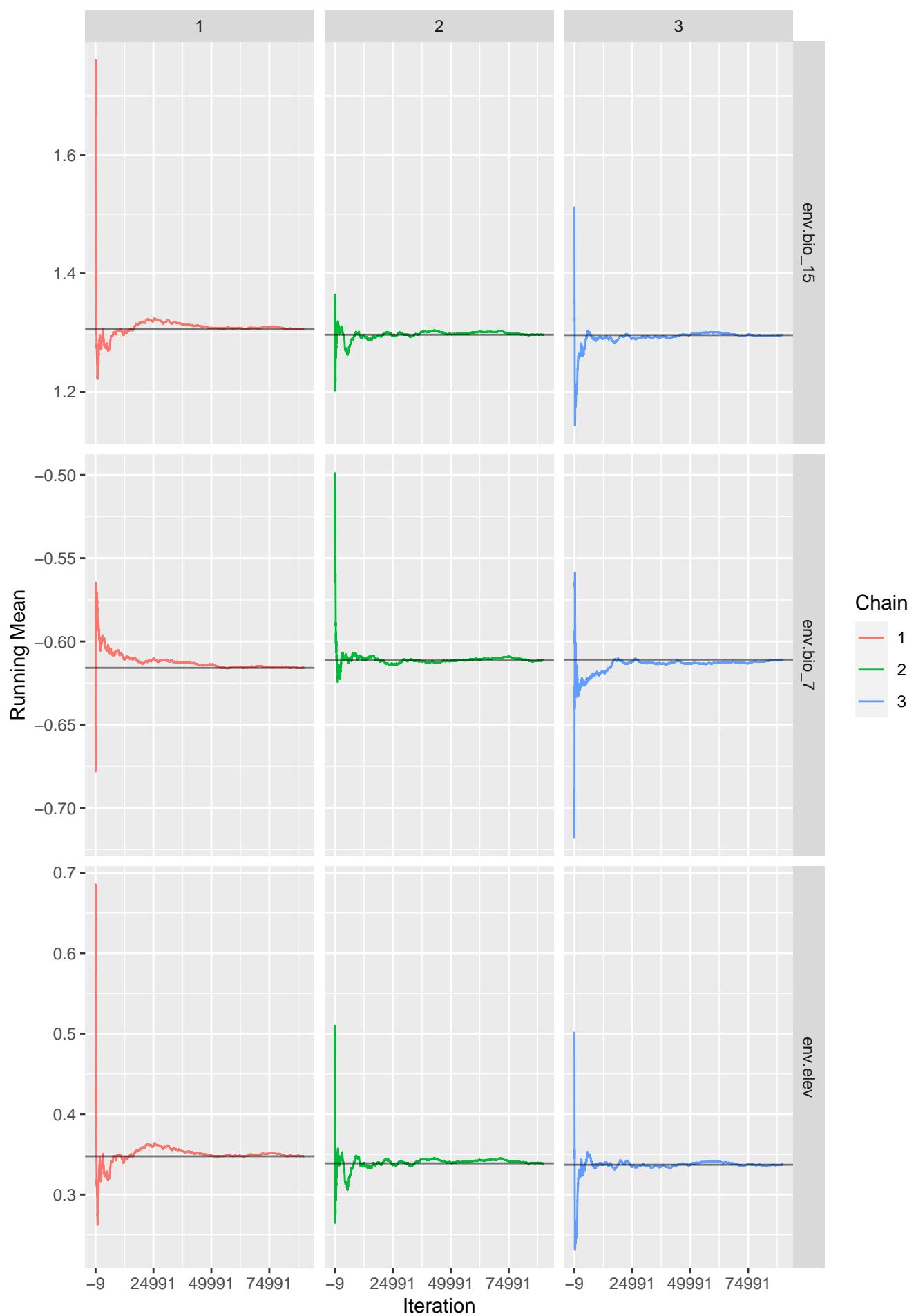
spline9

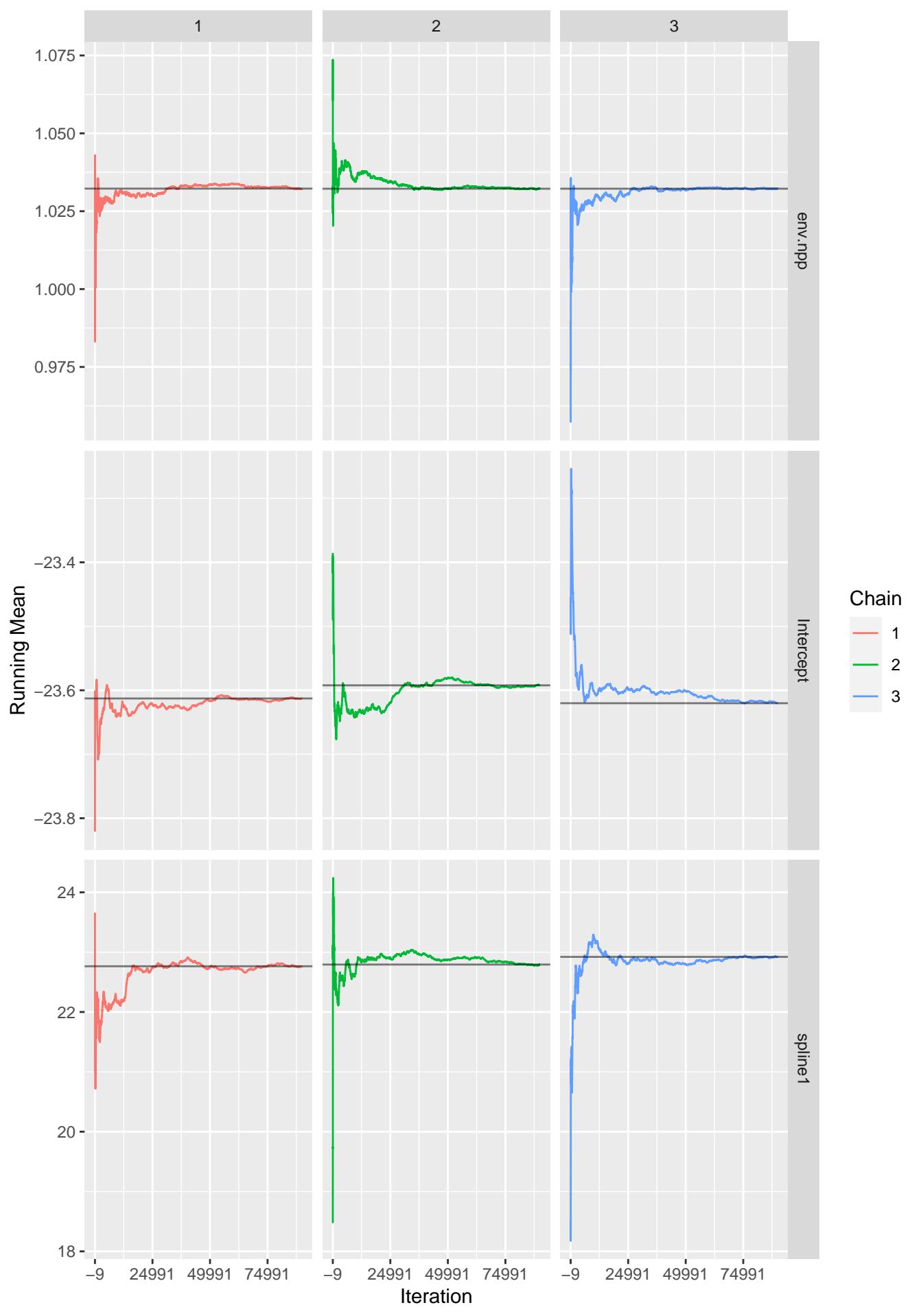


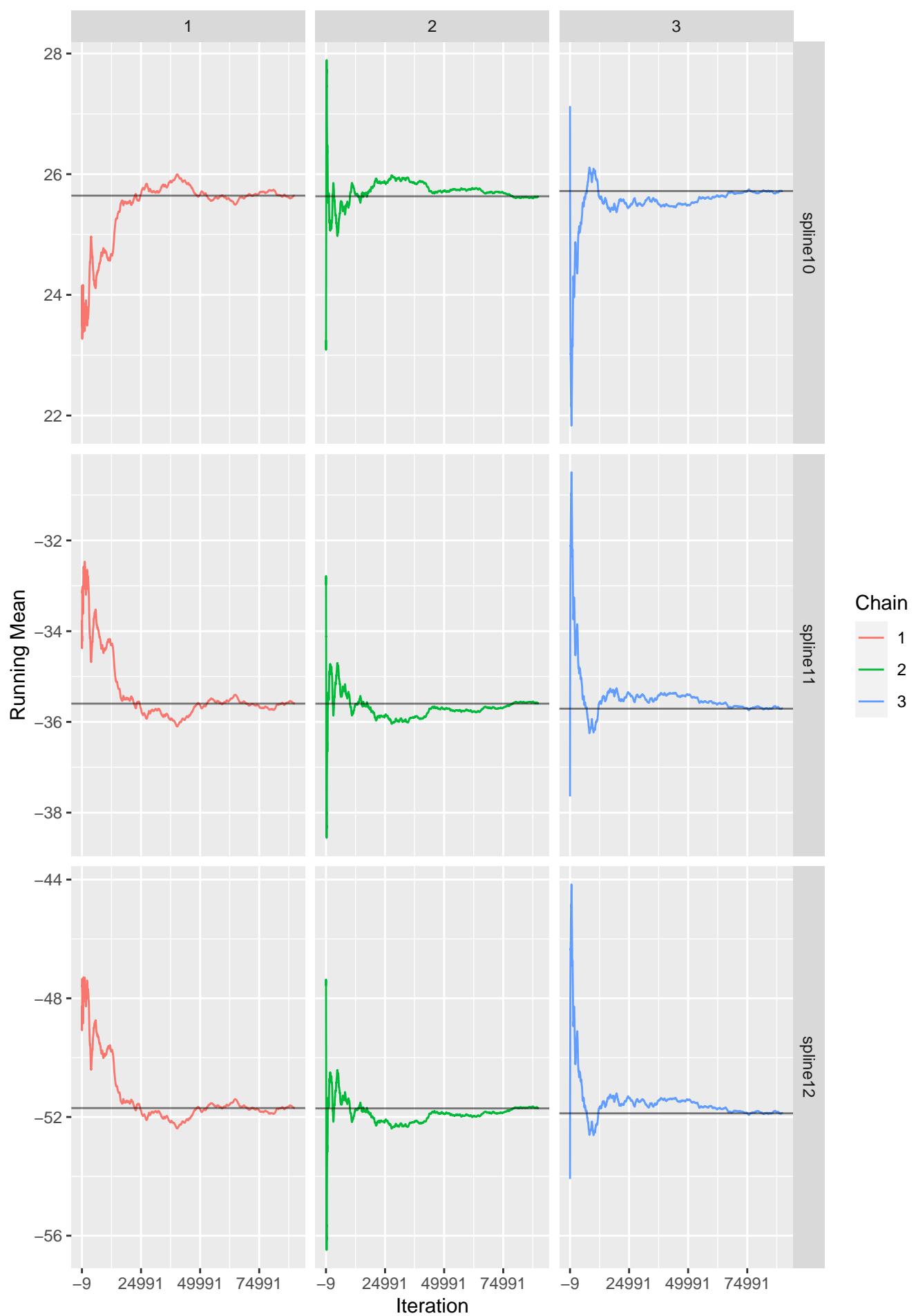
Chain

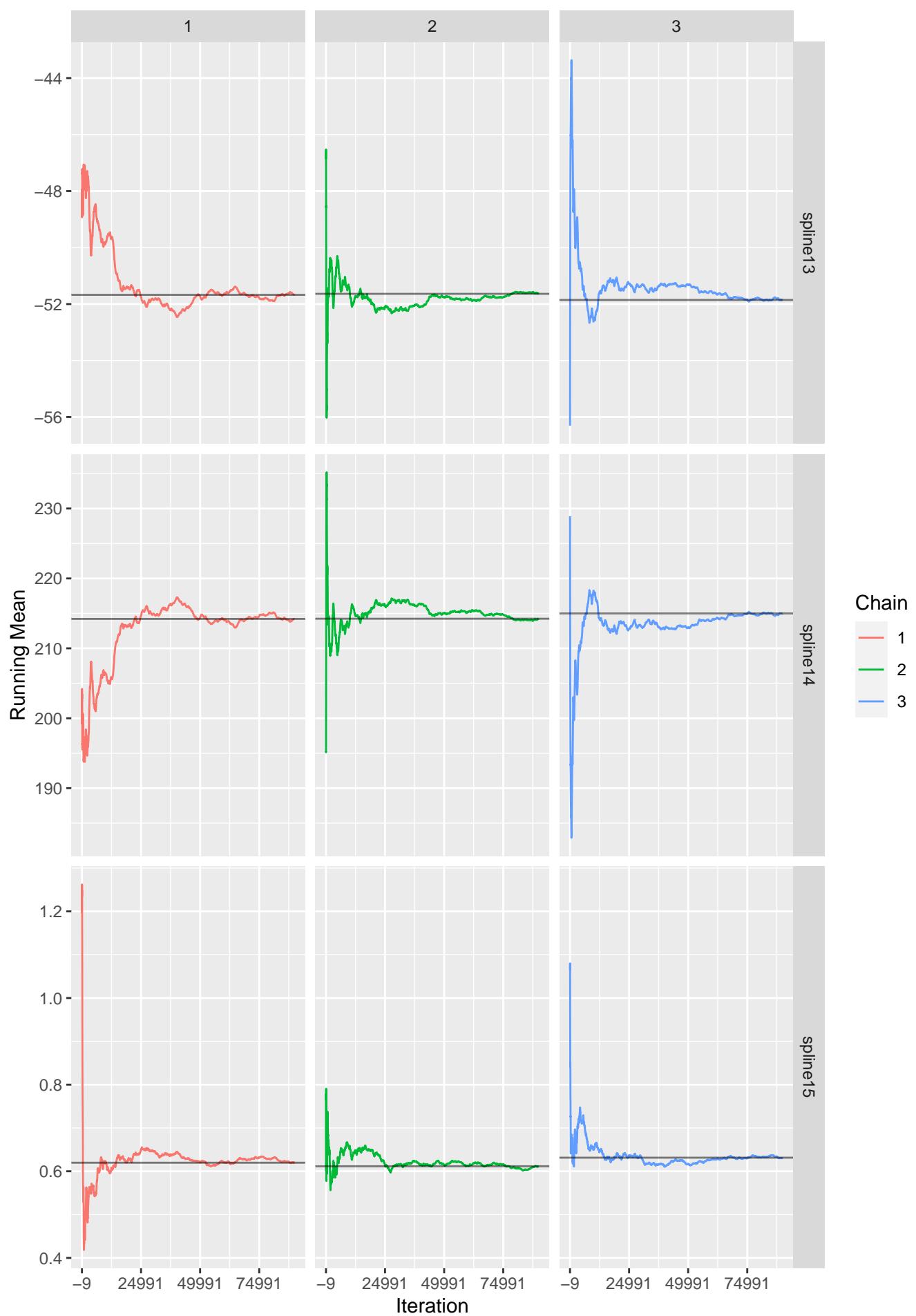
- 1
- 2
- 3

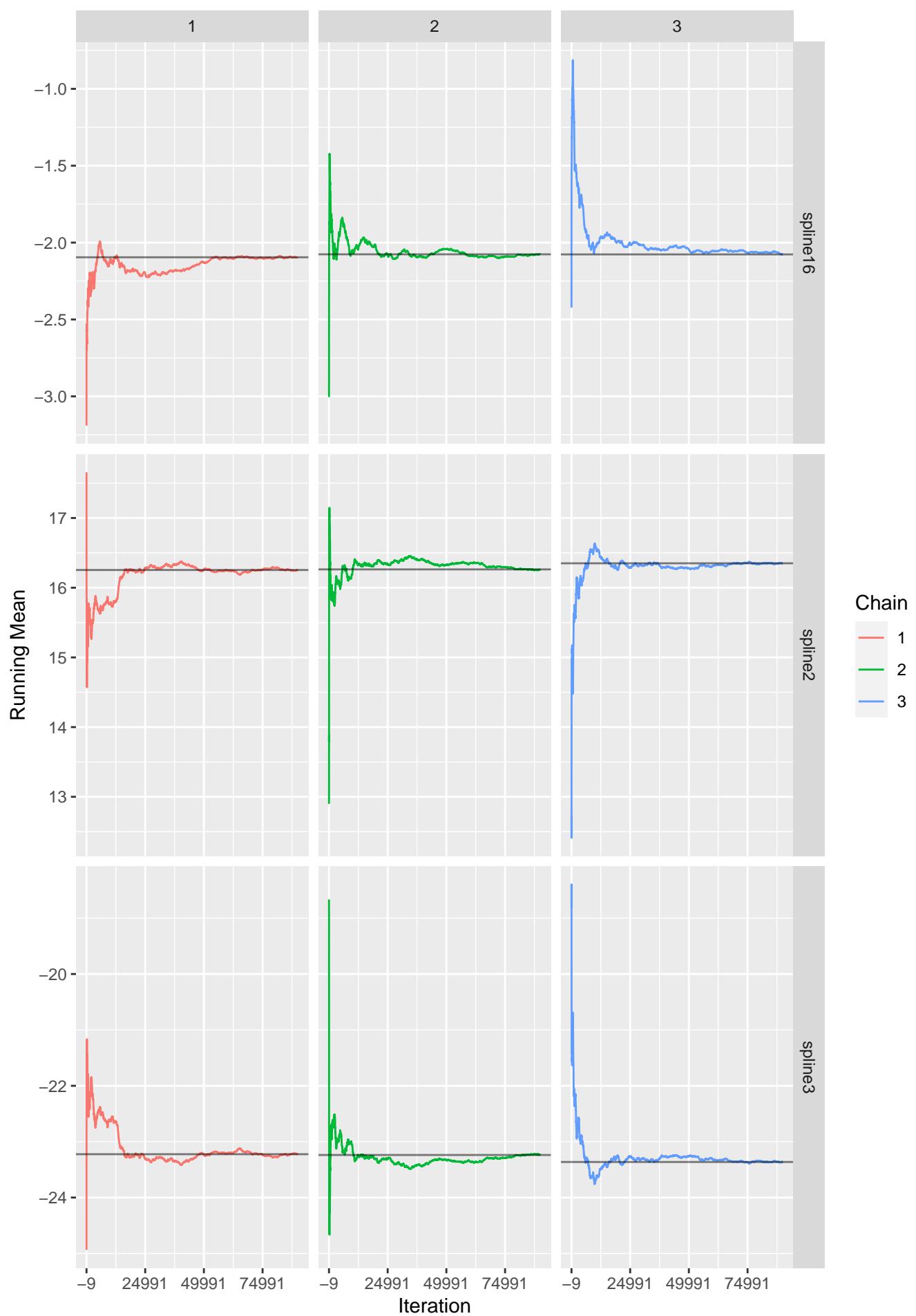
Iteration

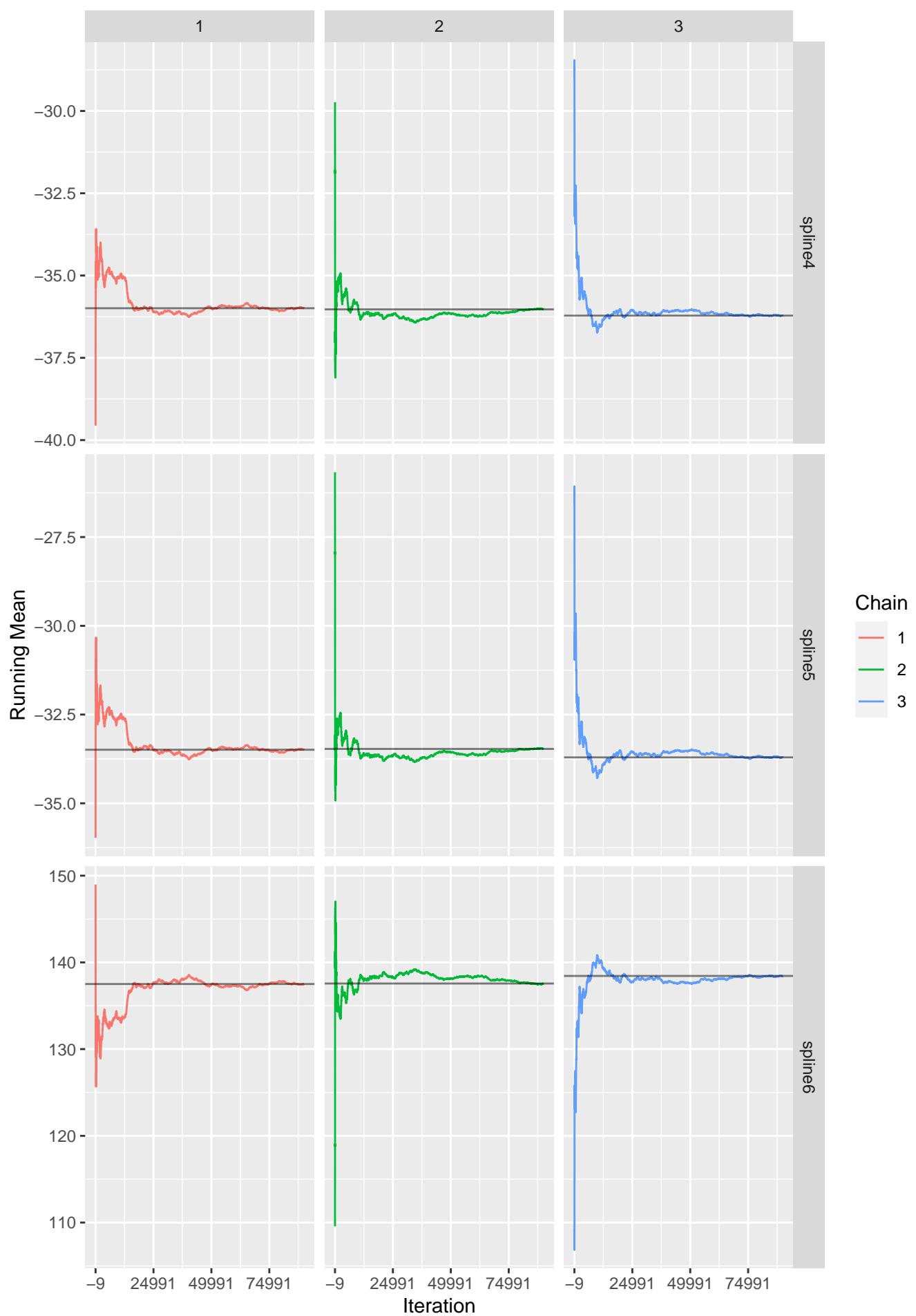


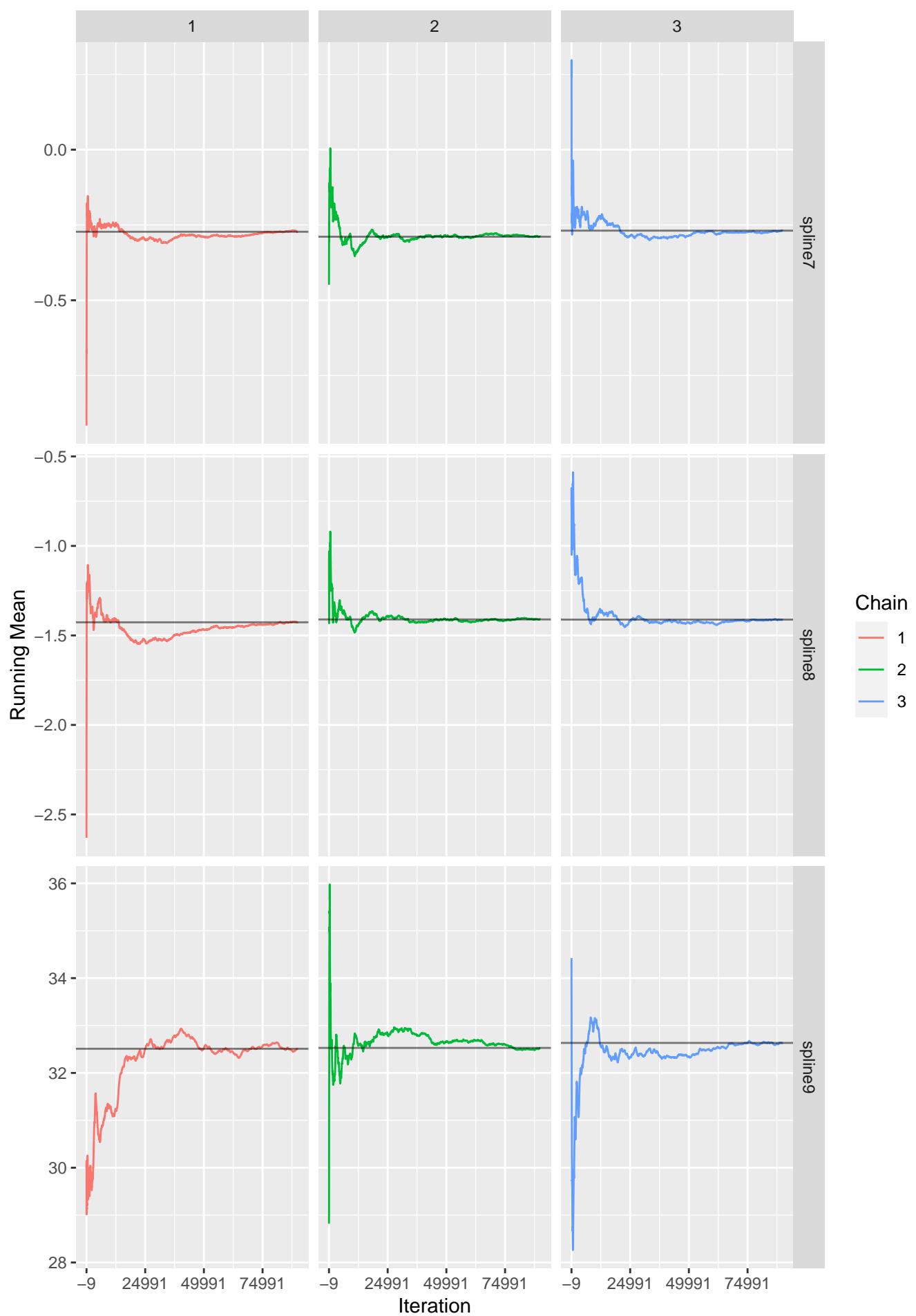


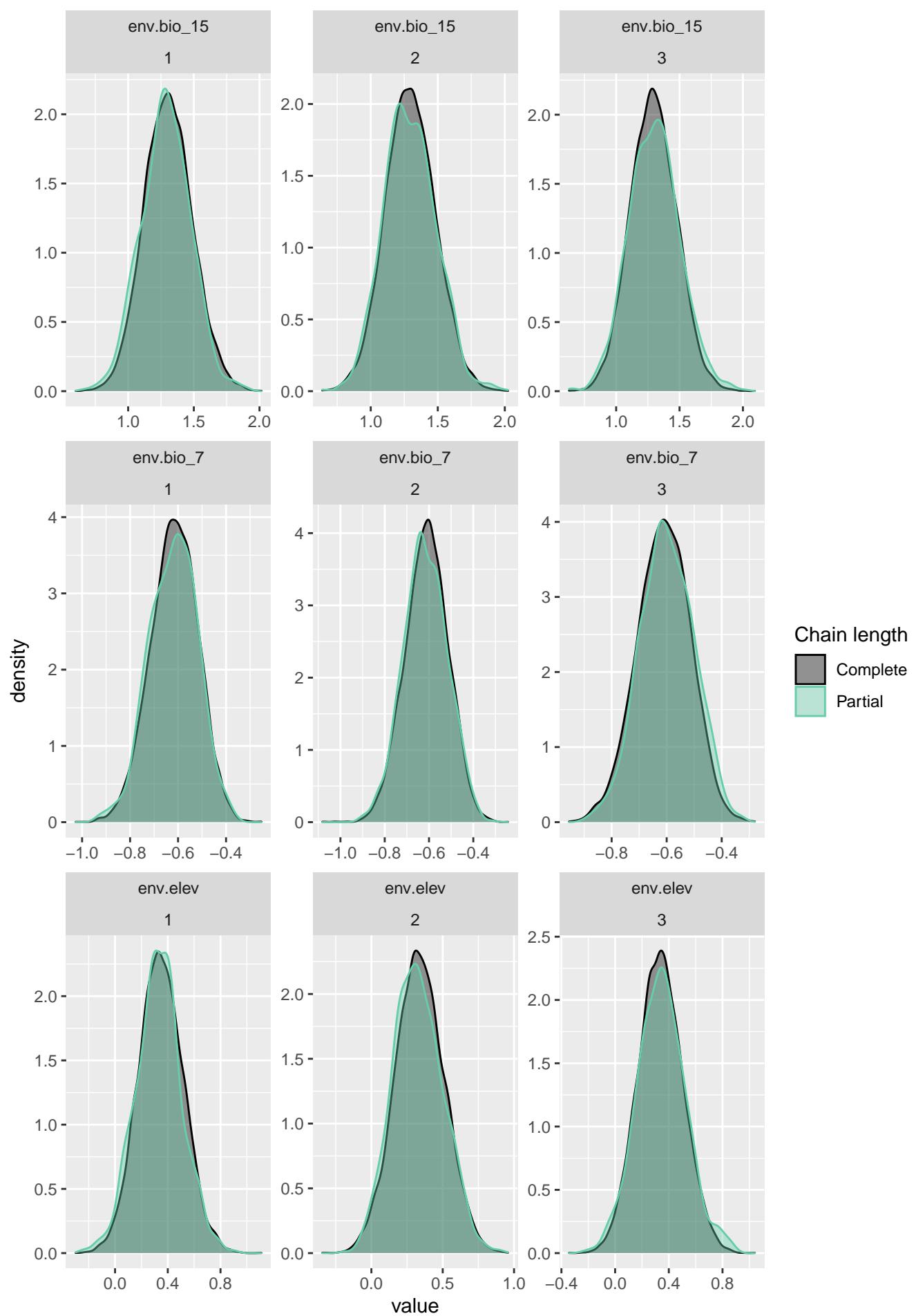


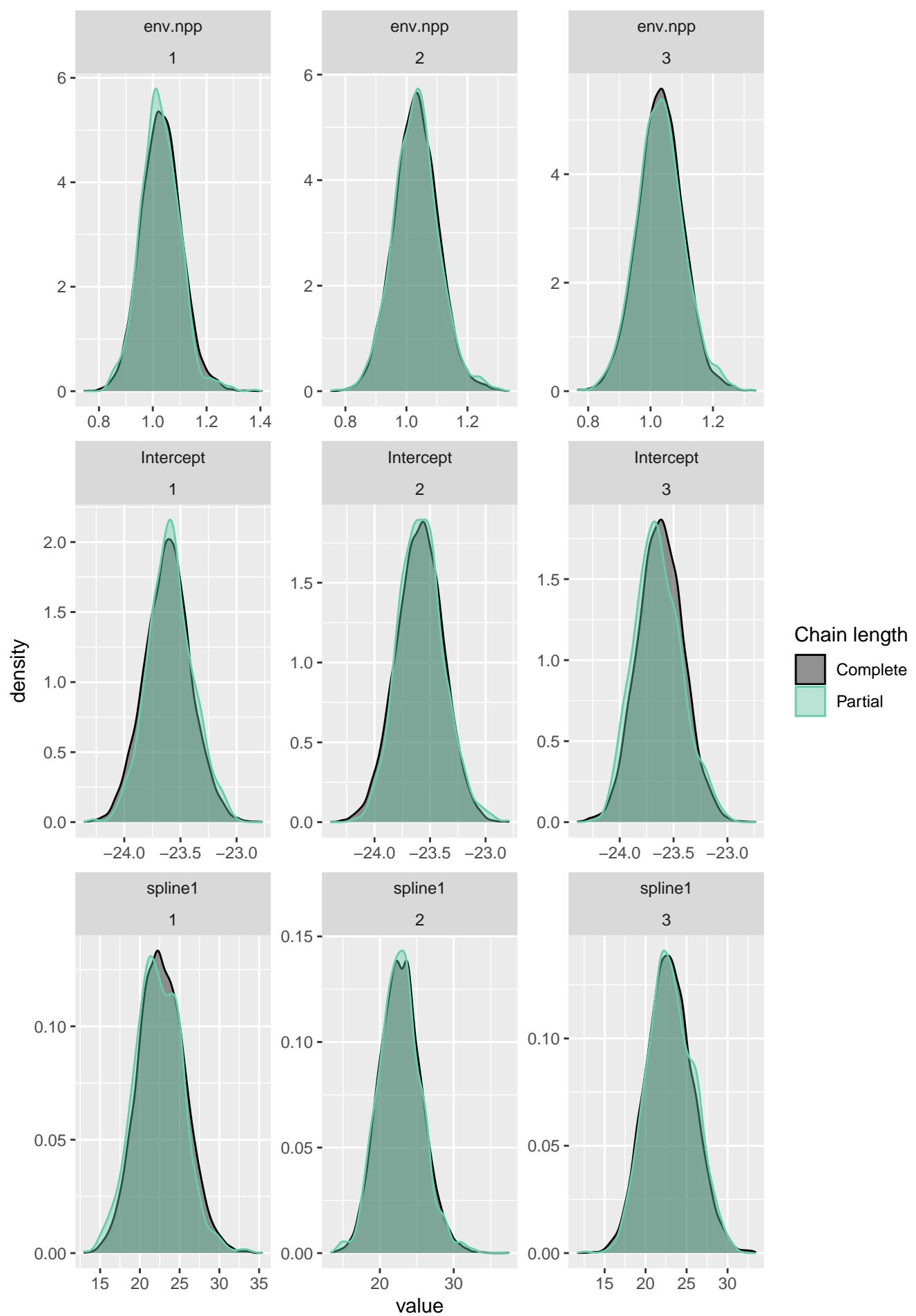


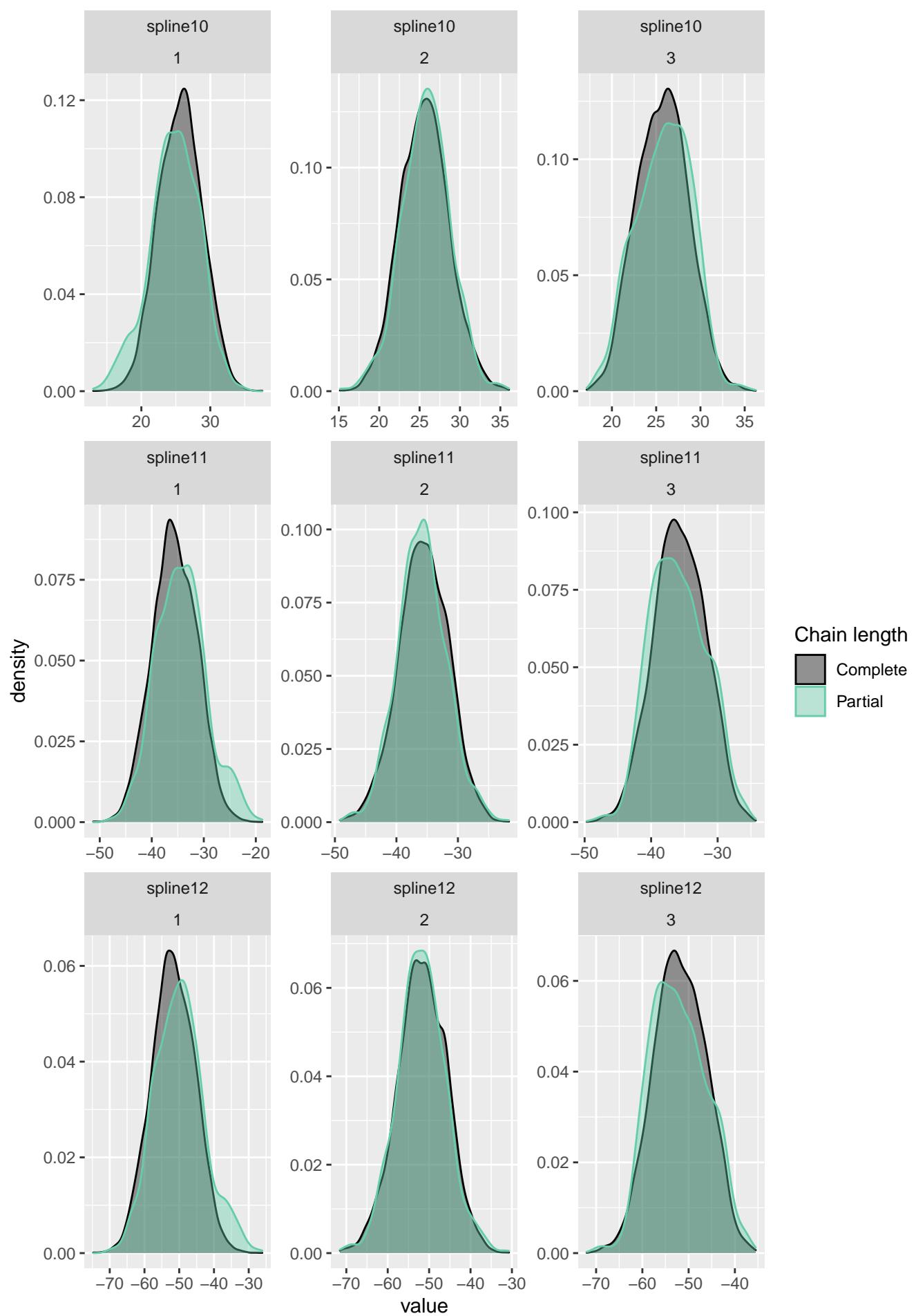


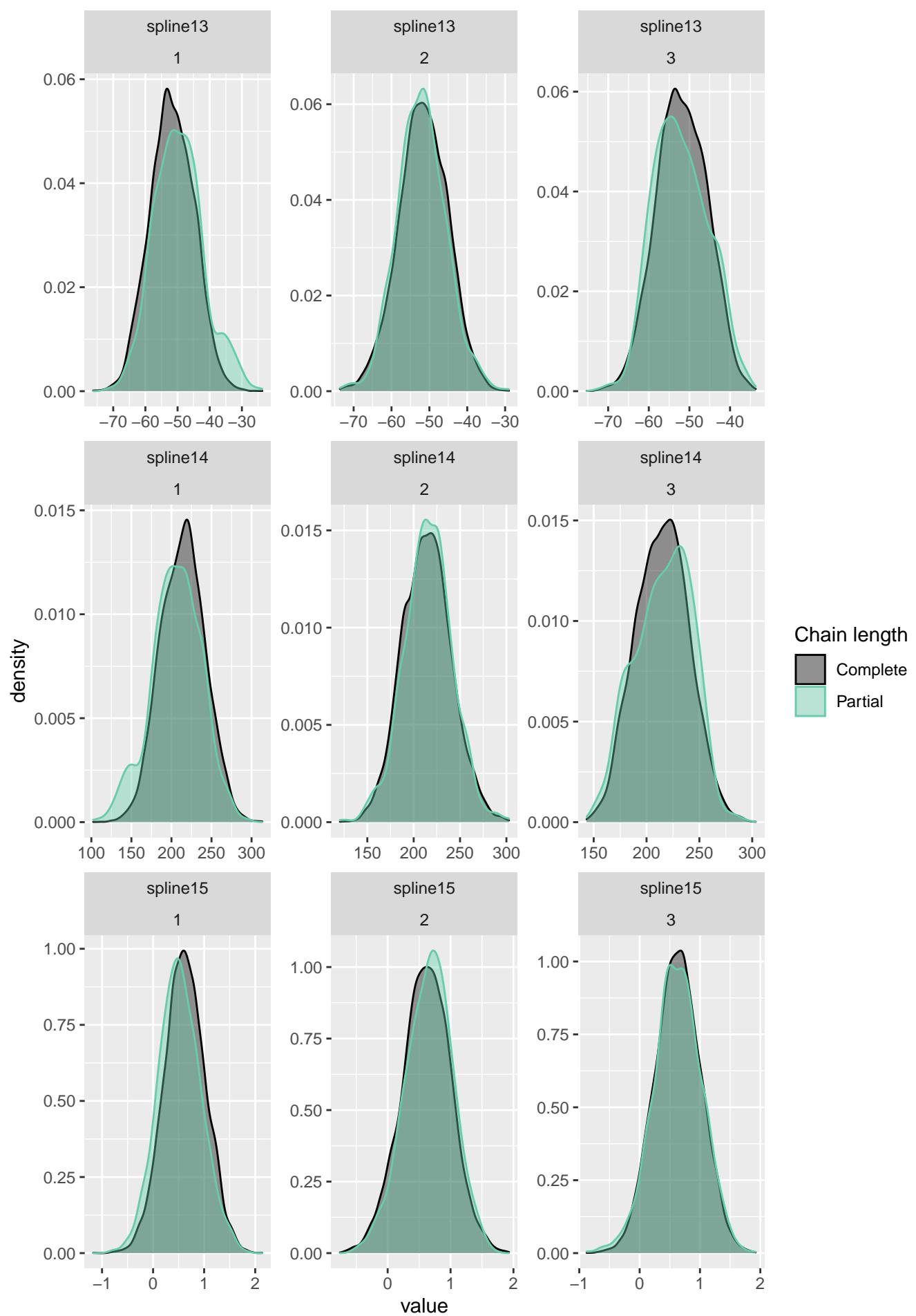


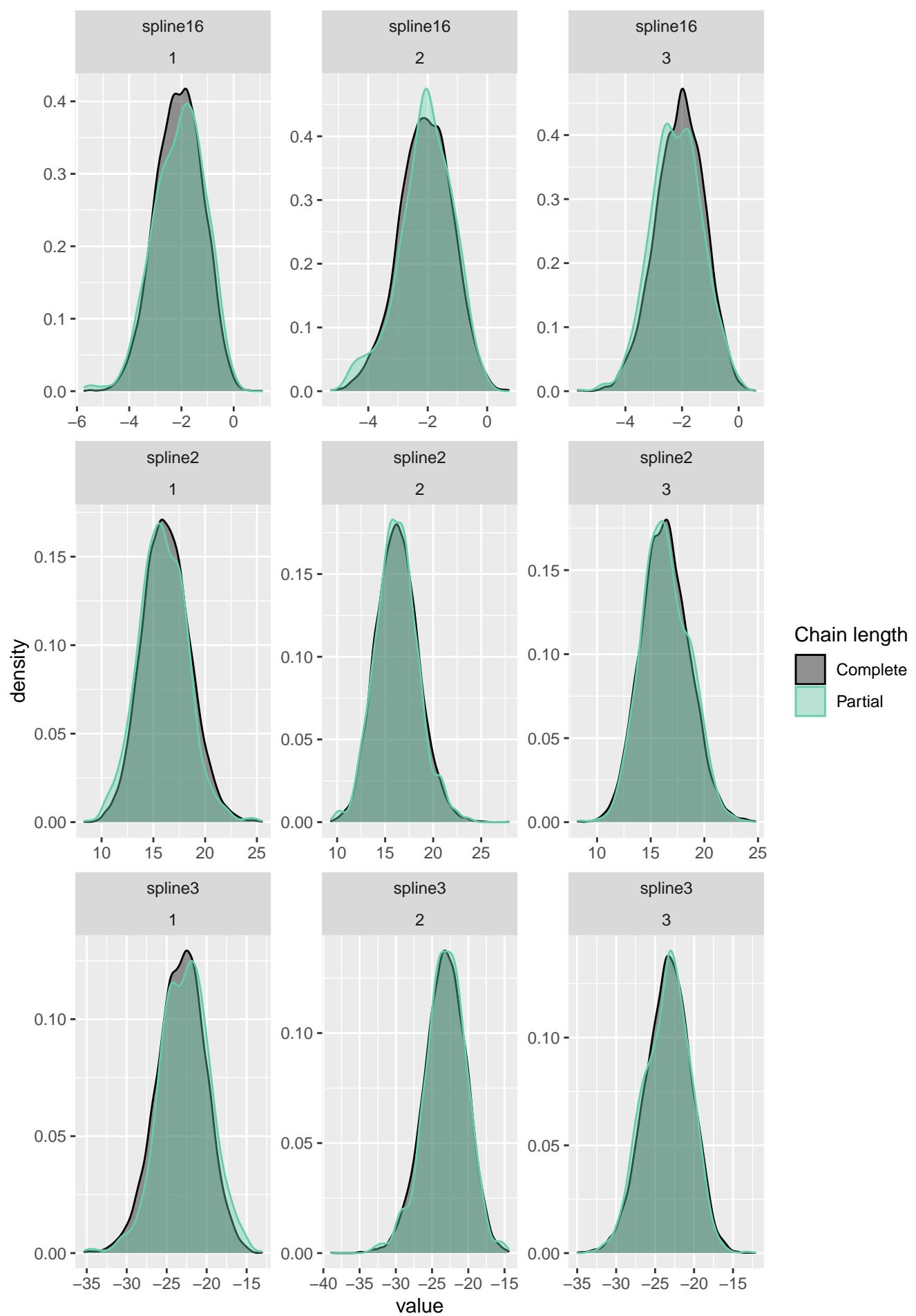


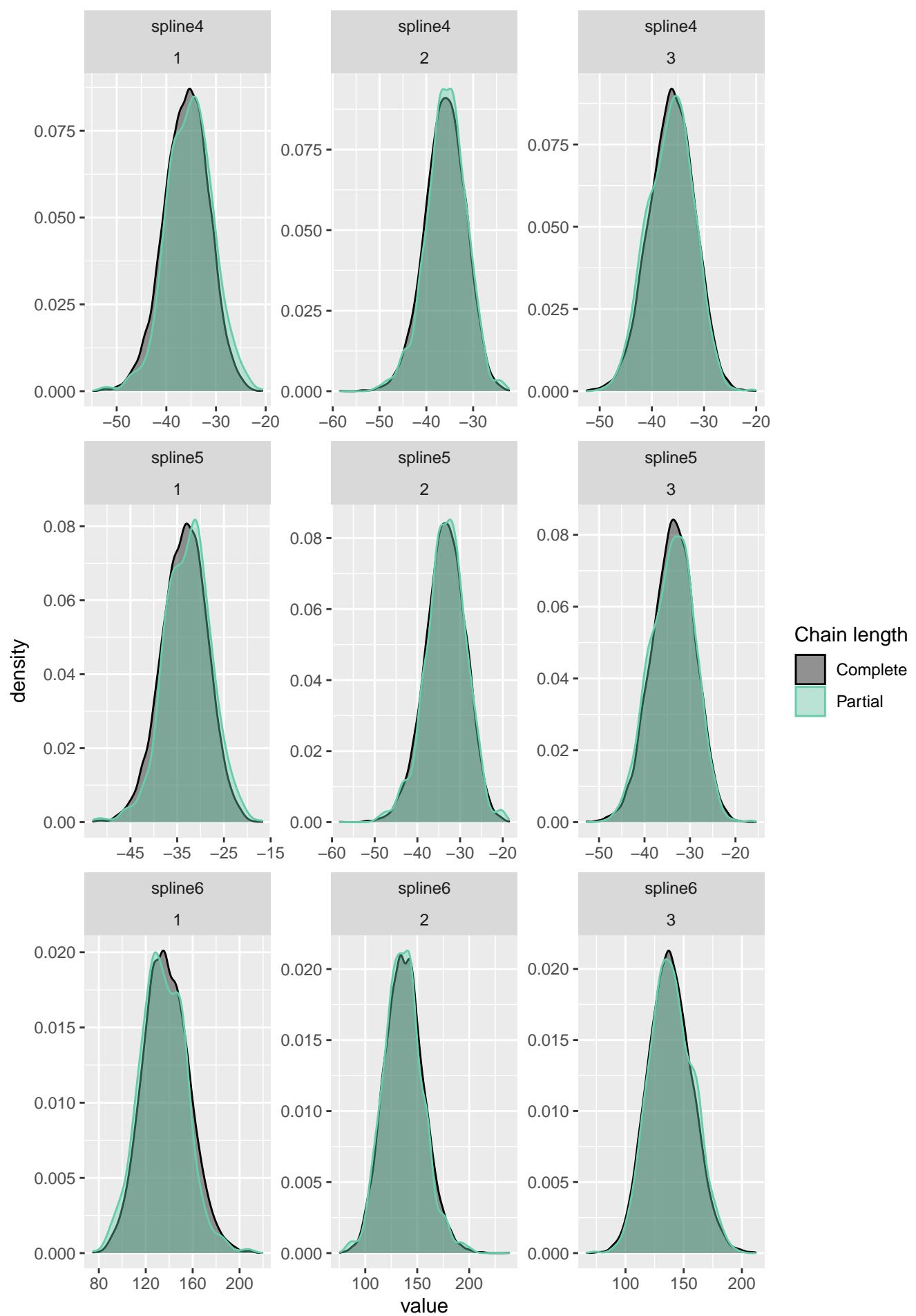


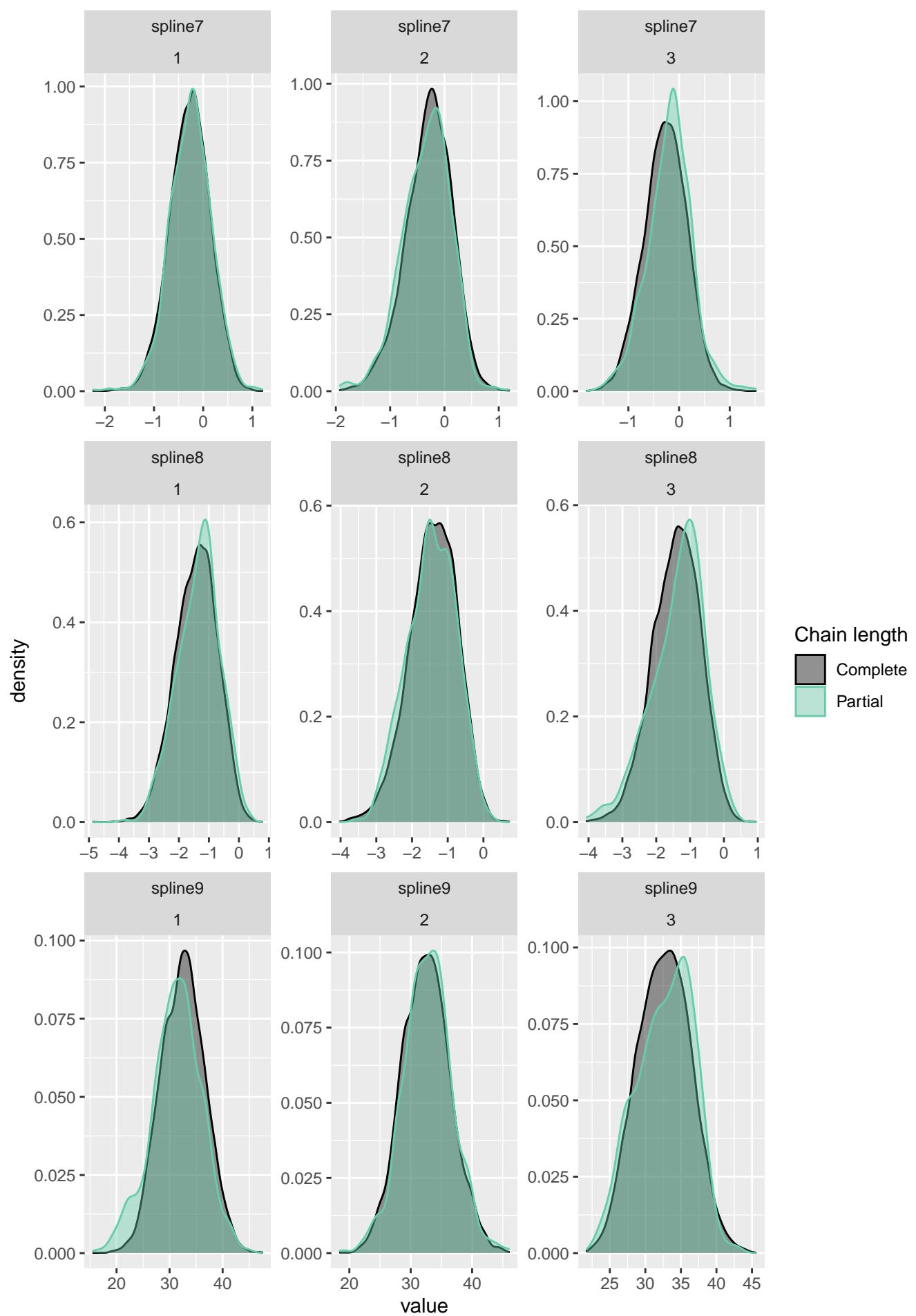


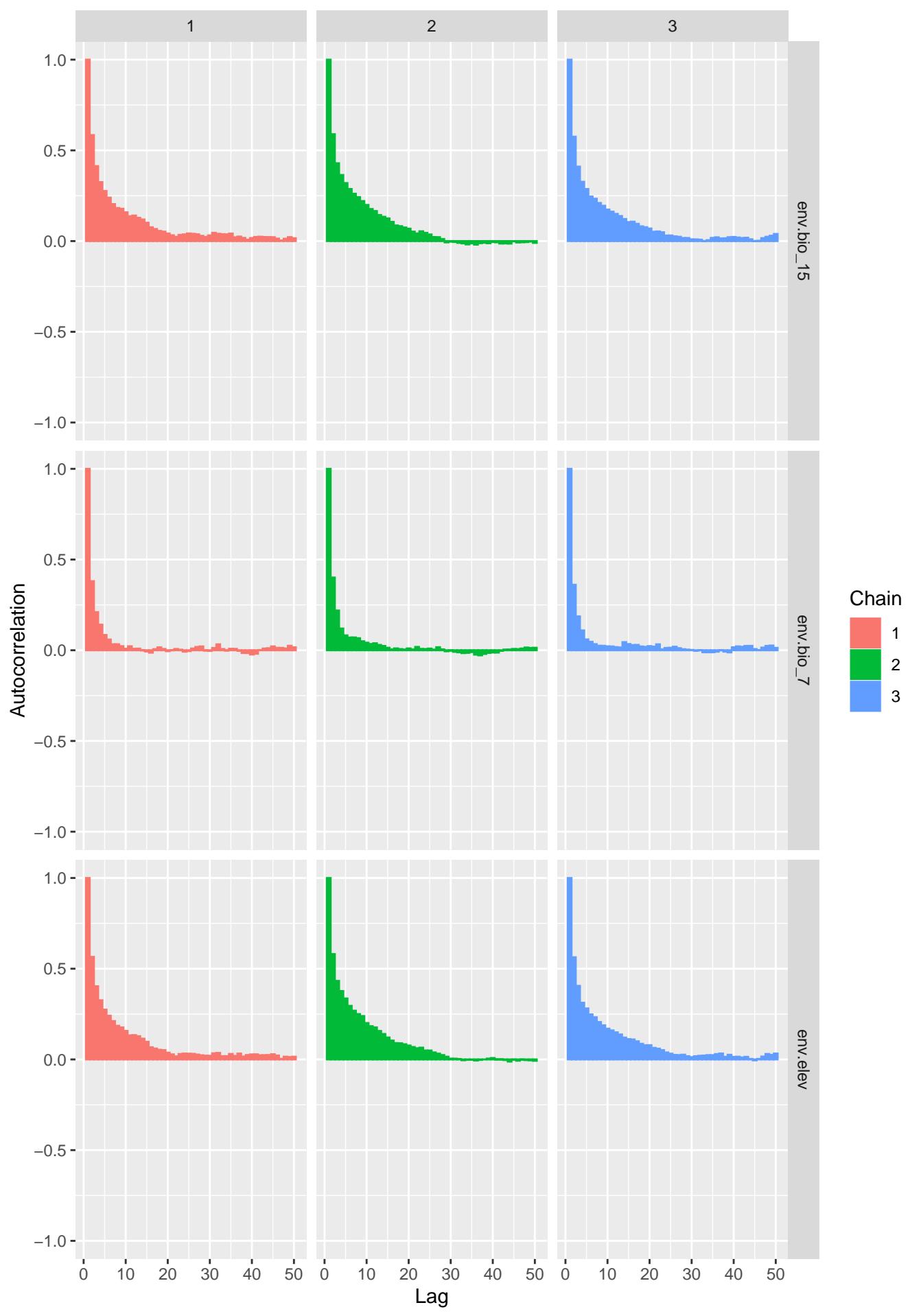


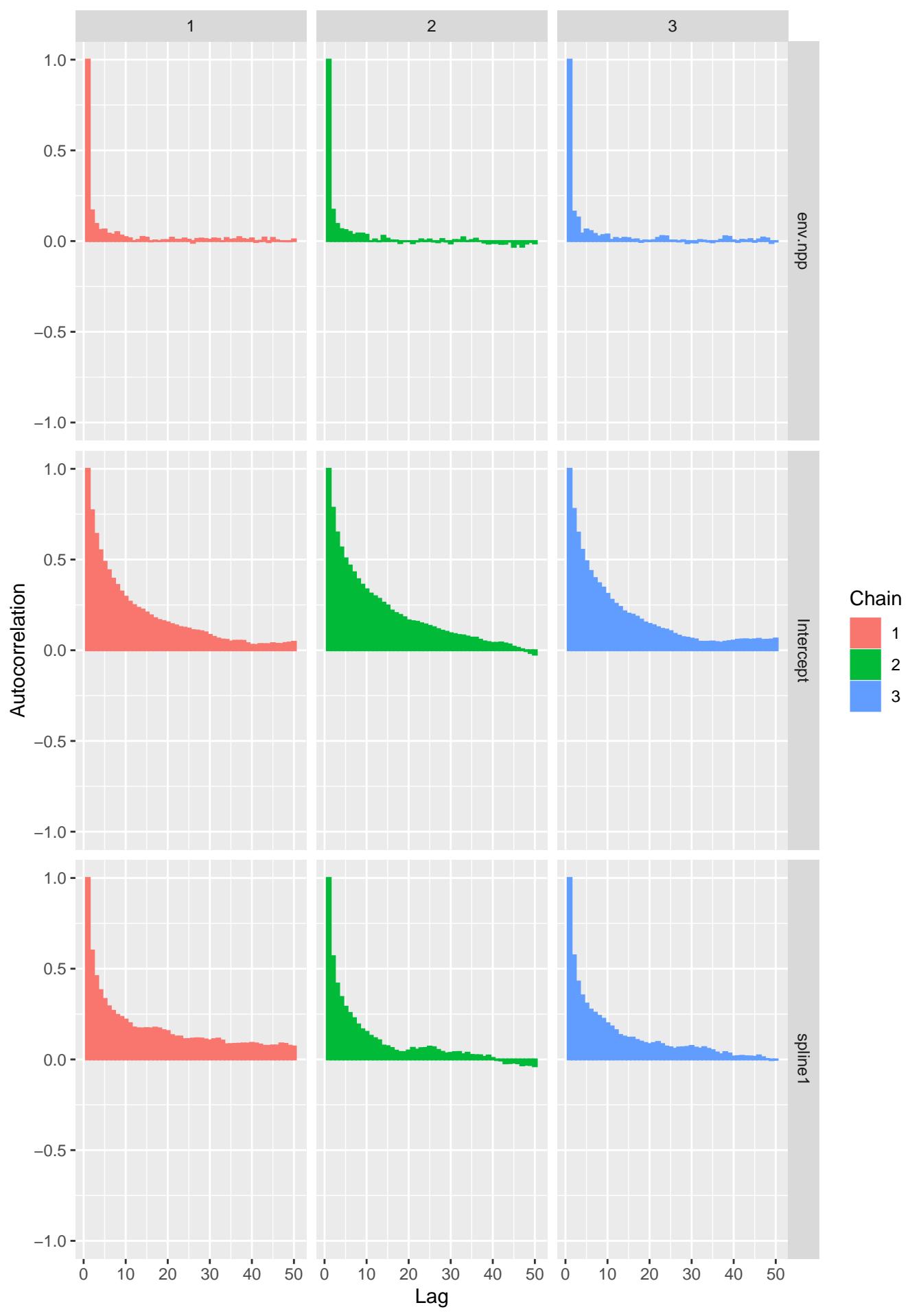


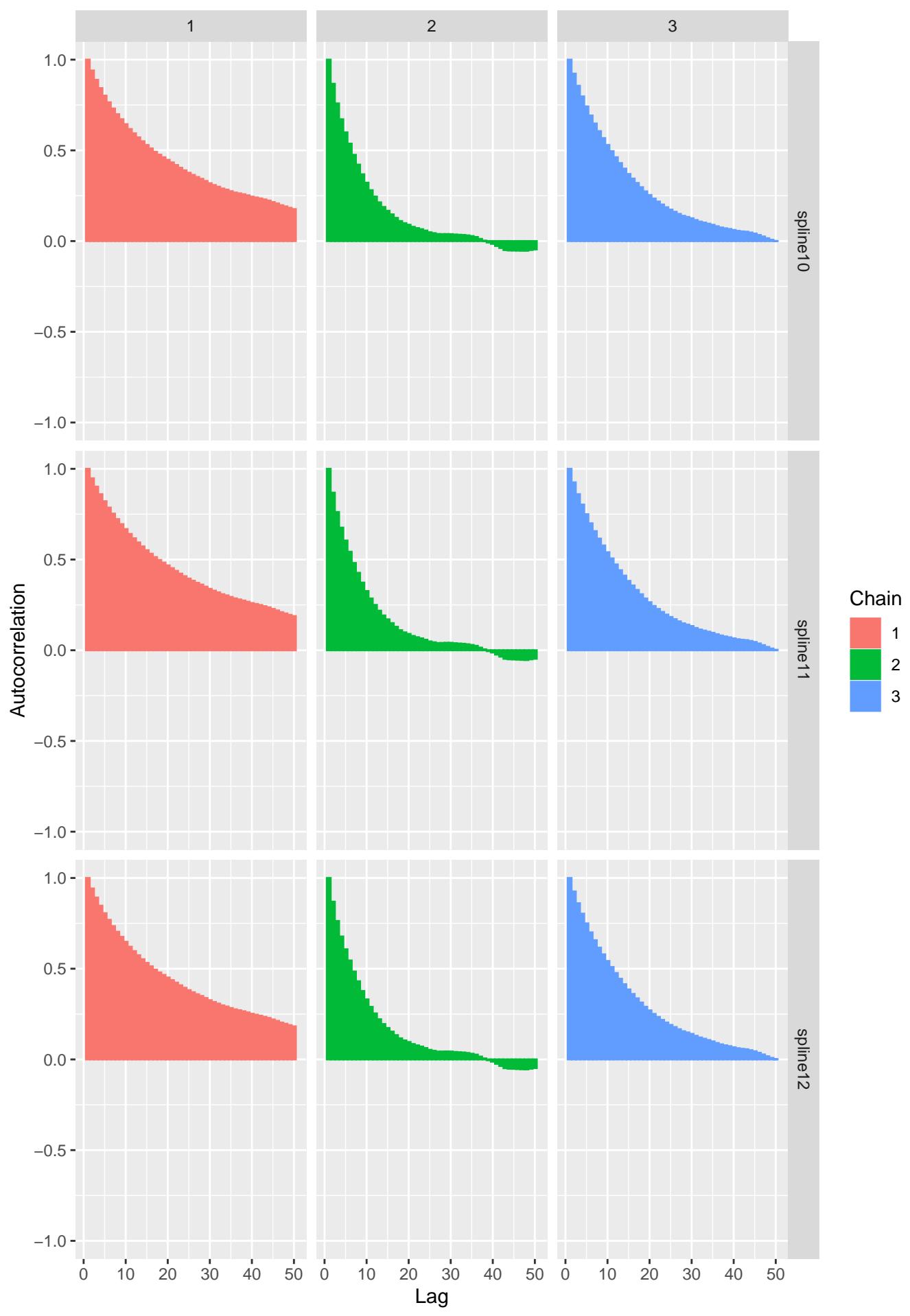


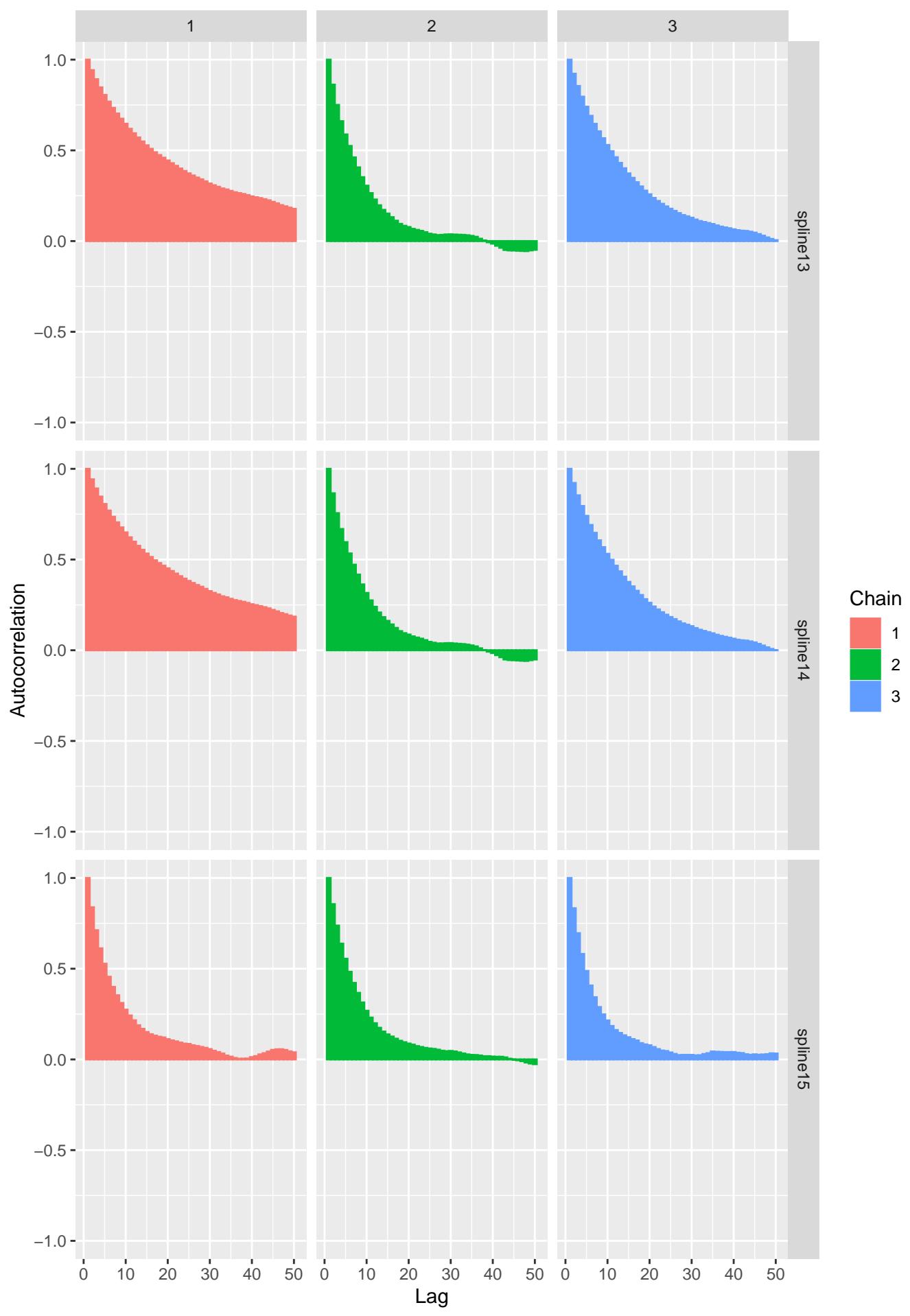


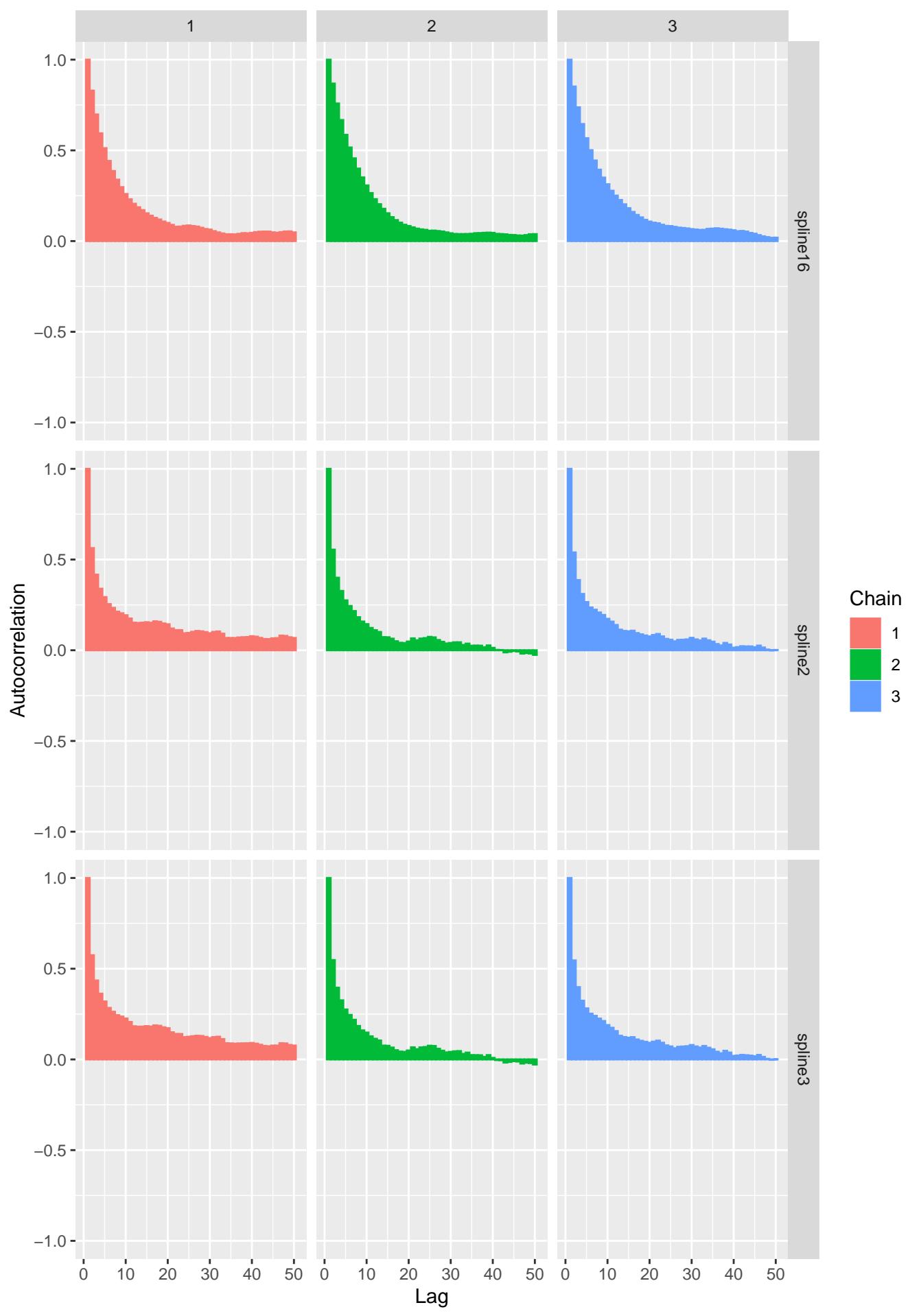


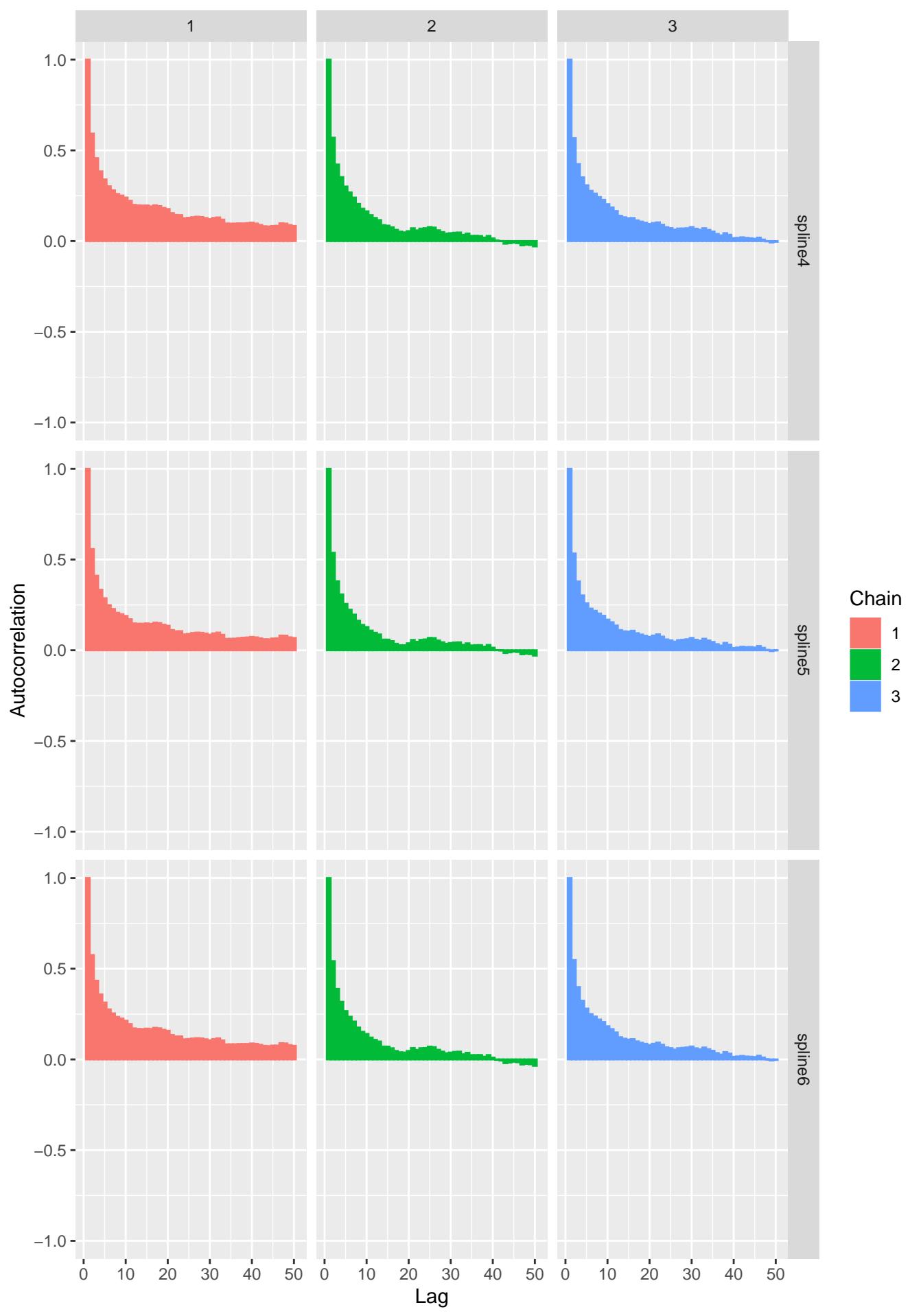


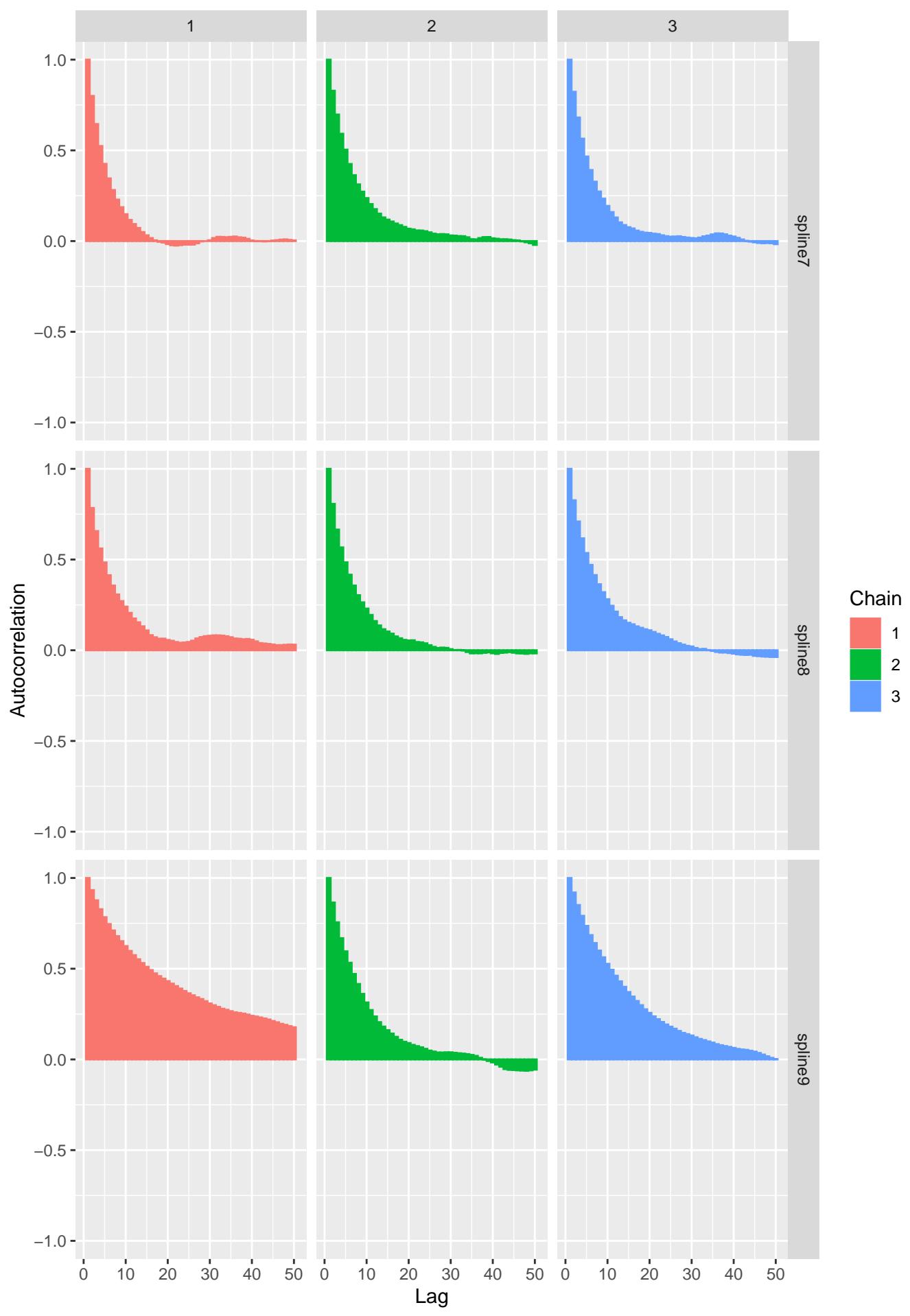


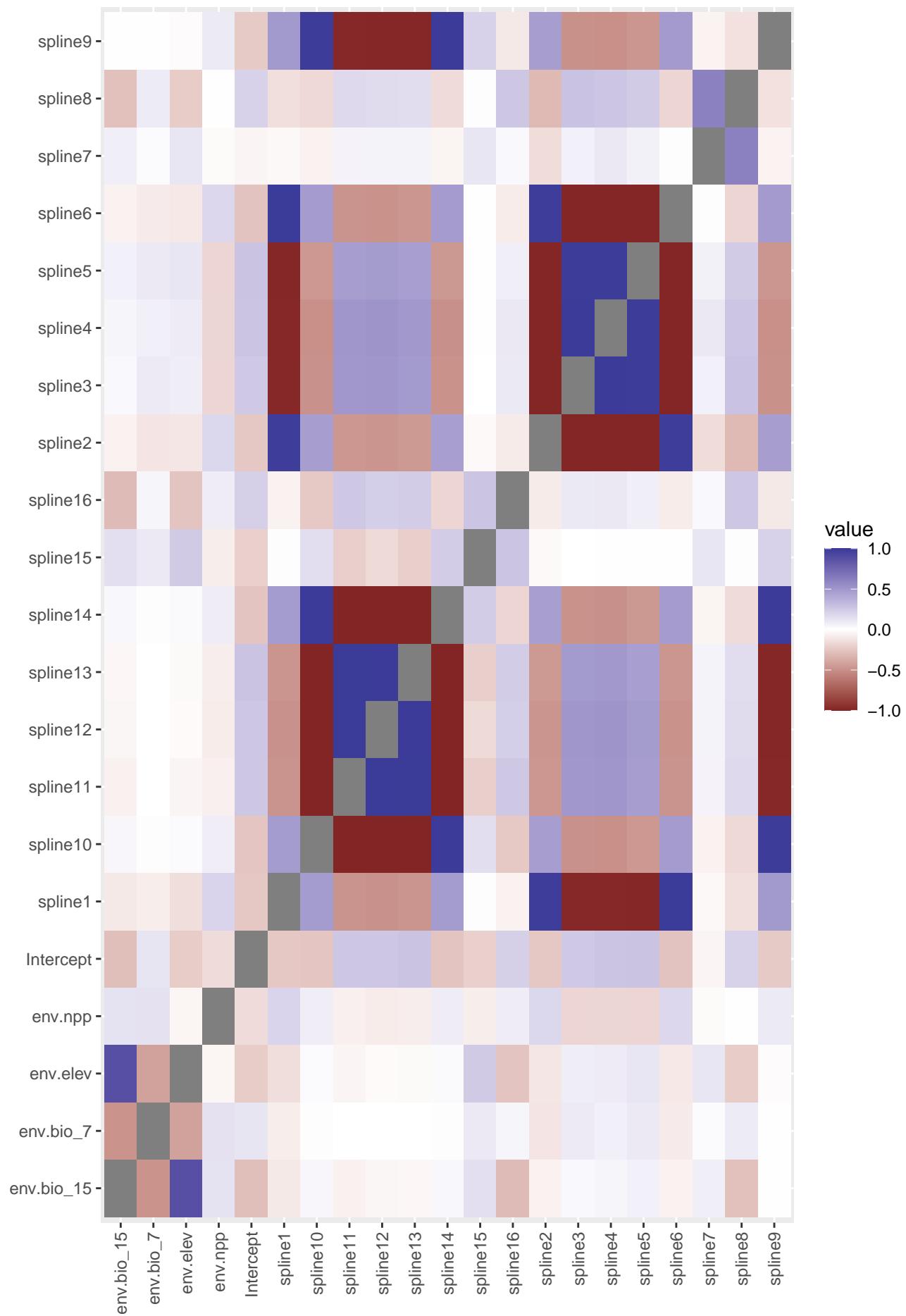




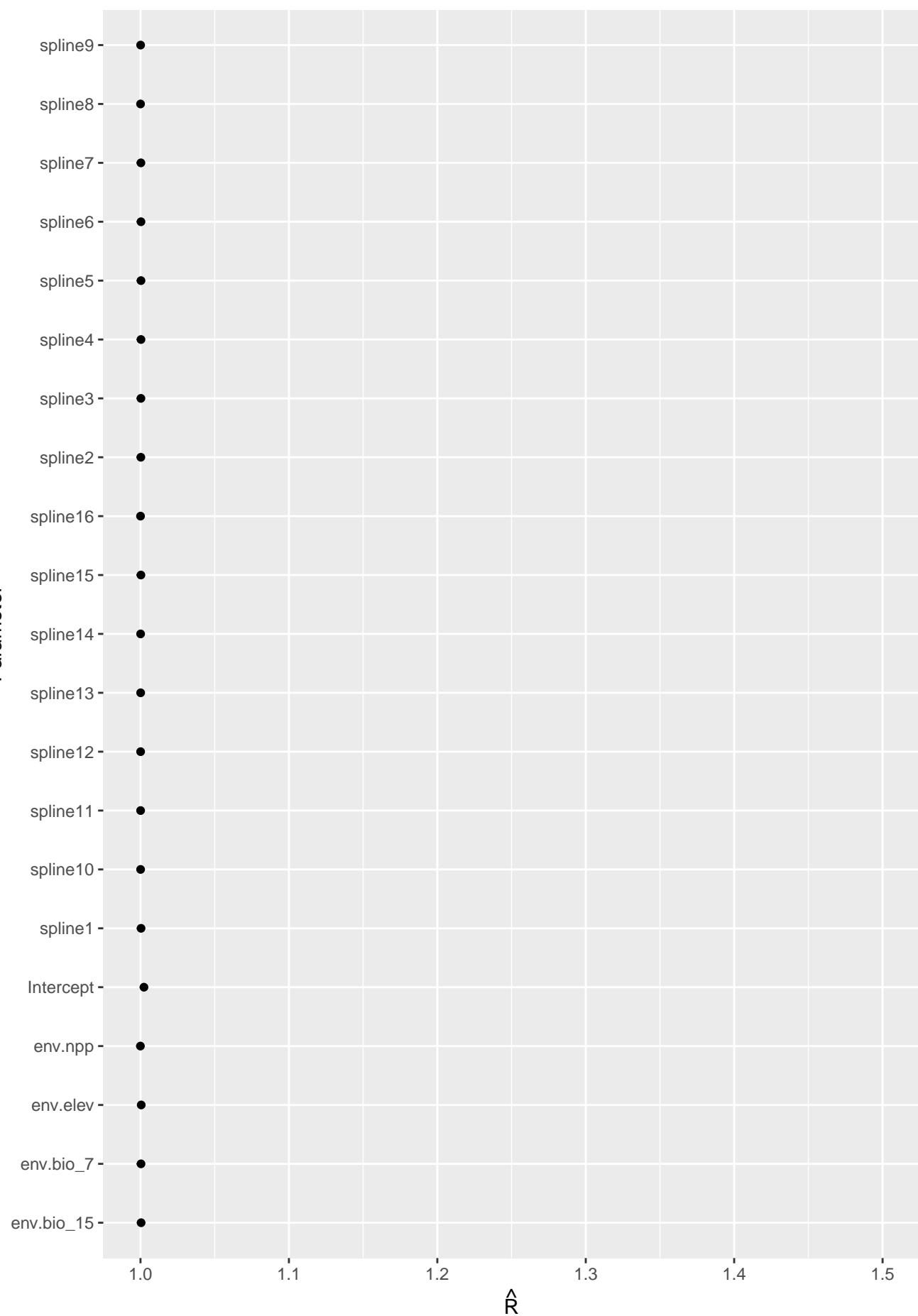




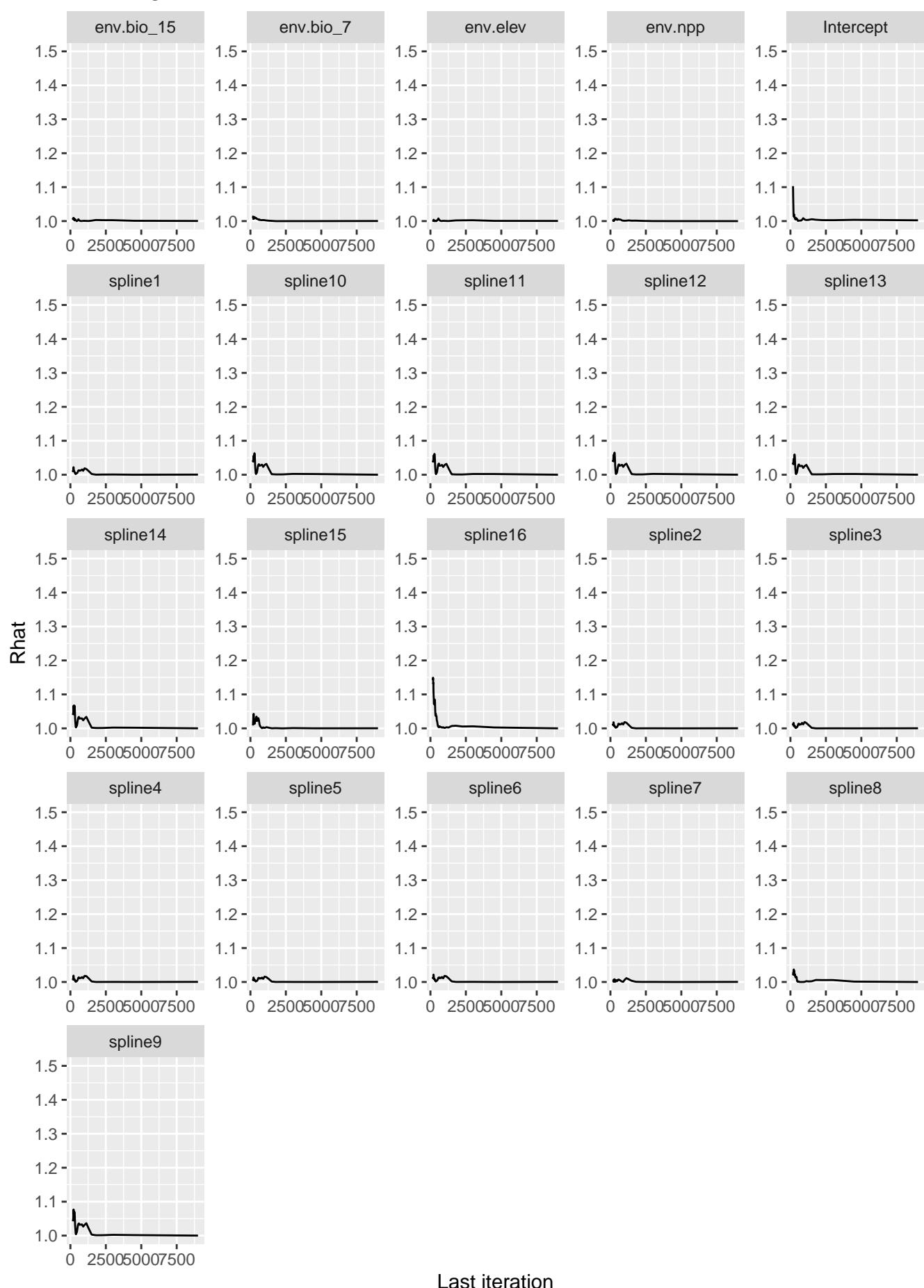




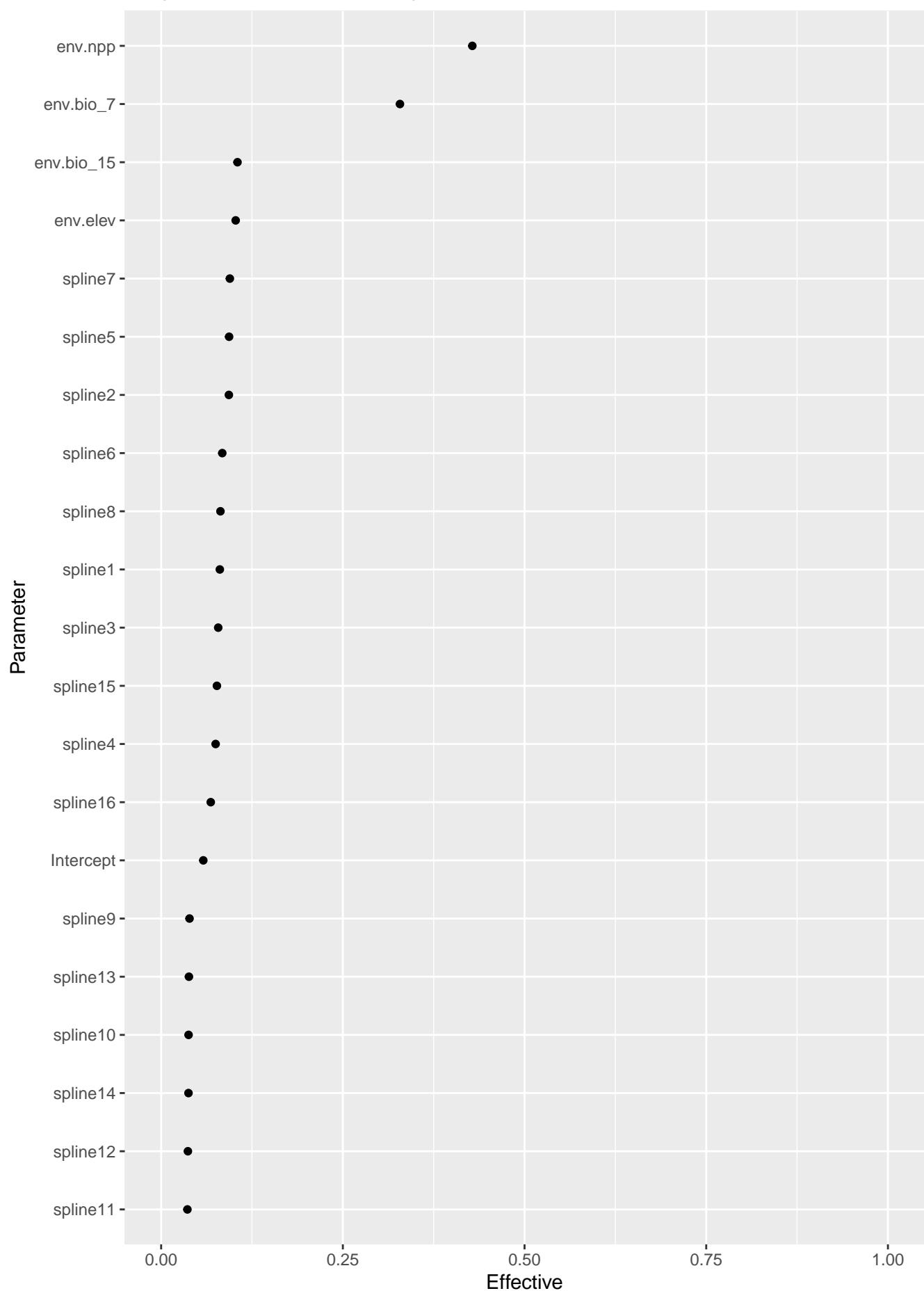
Potential Scale Reduction Factors



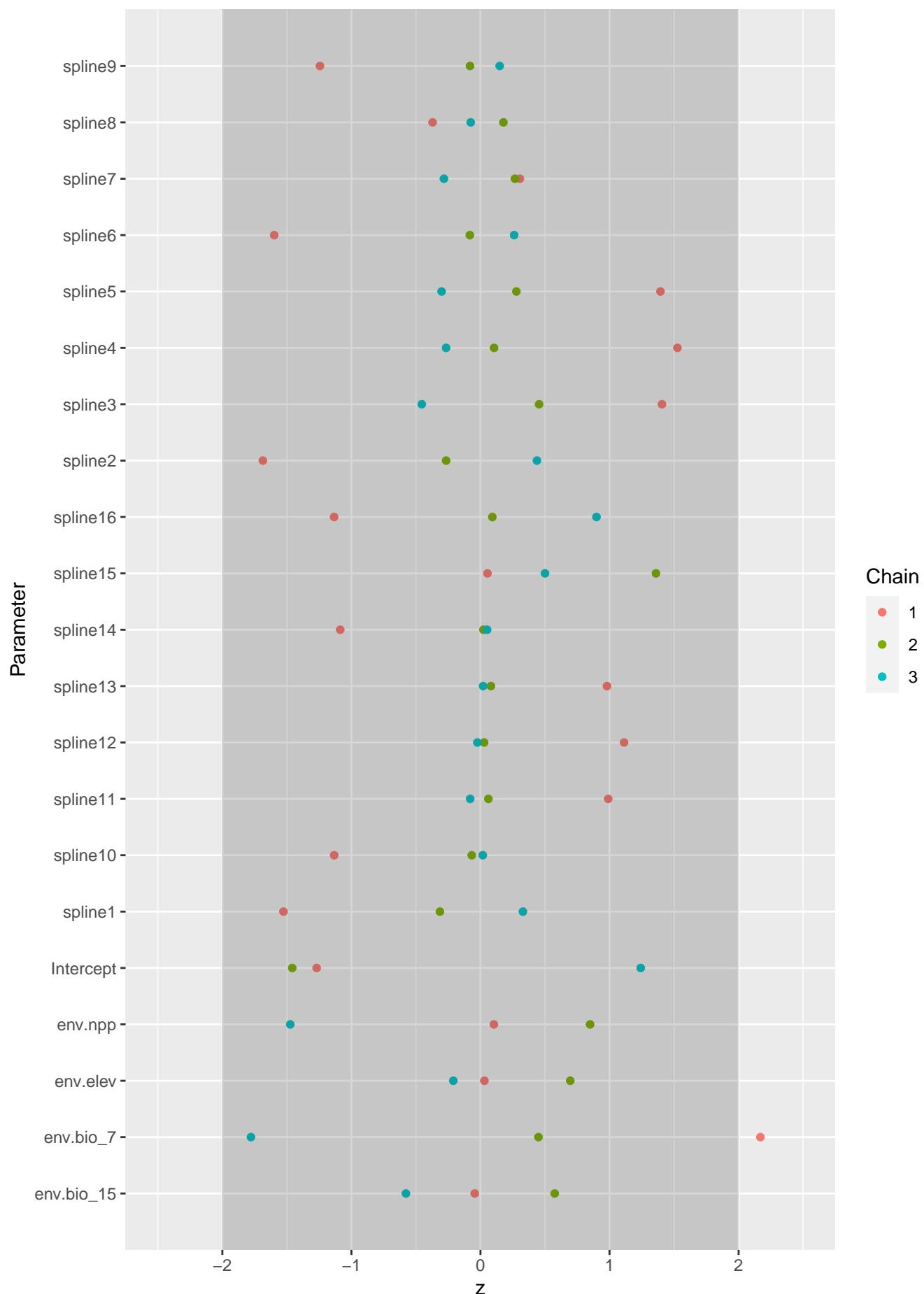
Shrinkage of Potential Scale Reduction Factors



Proportion of effective independent draws



Geweke Diagnostics



b