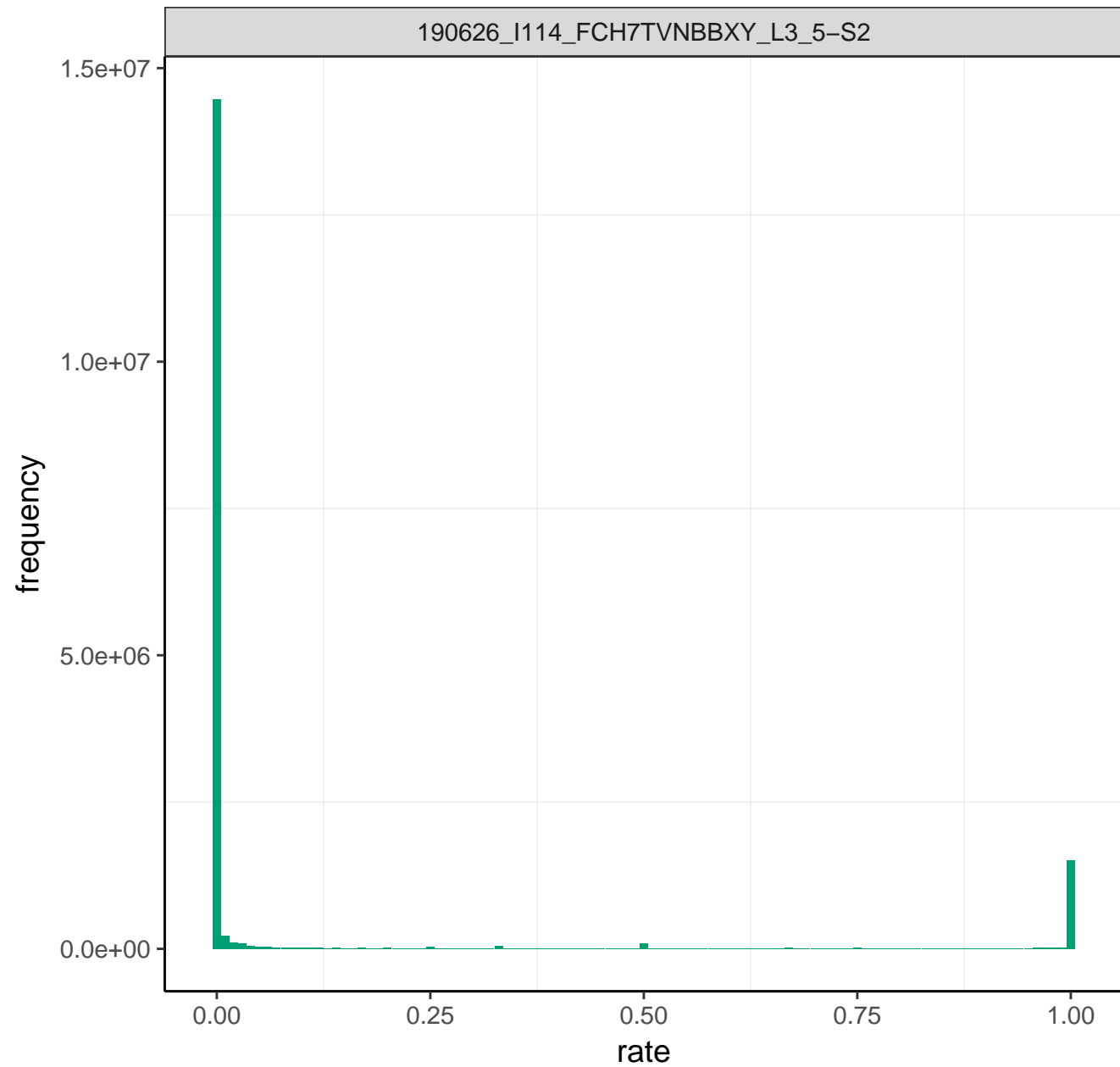


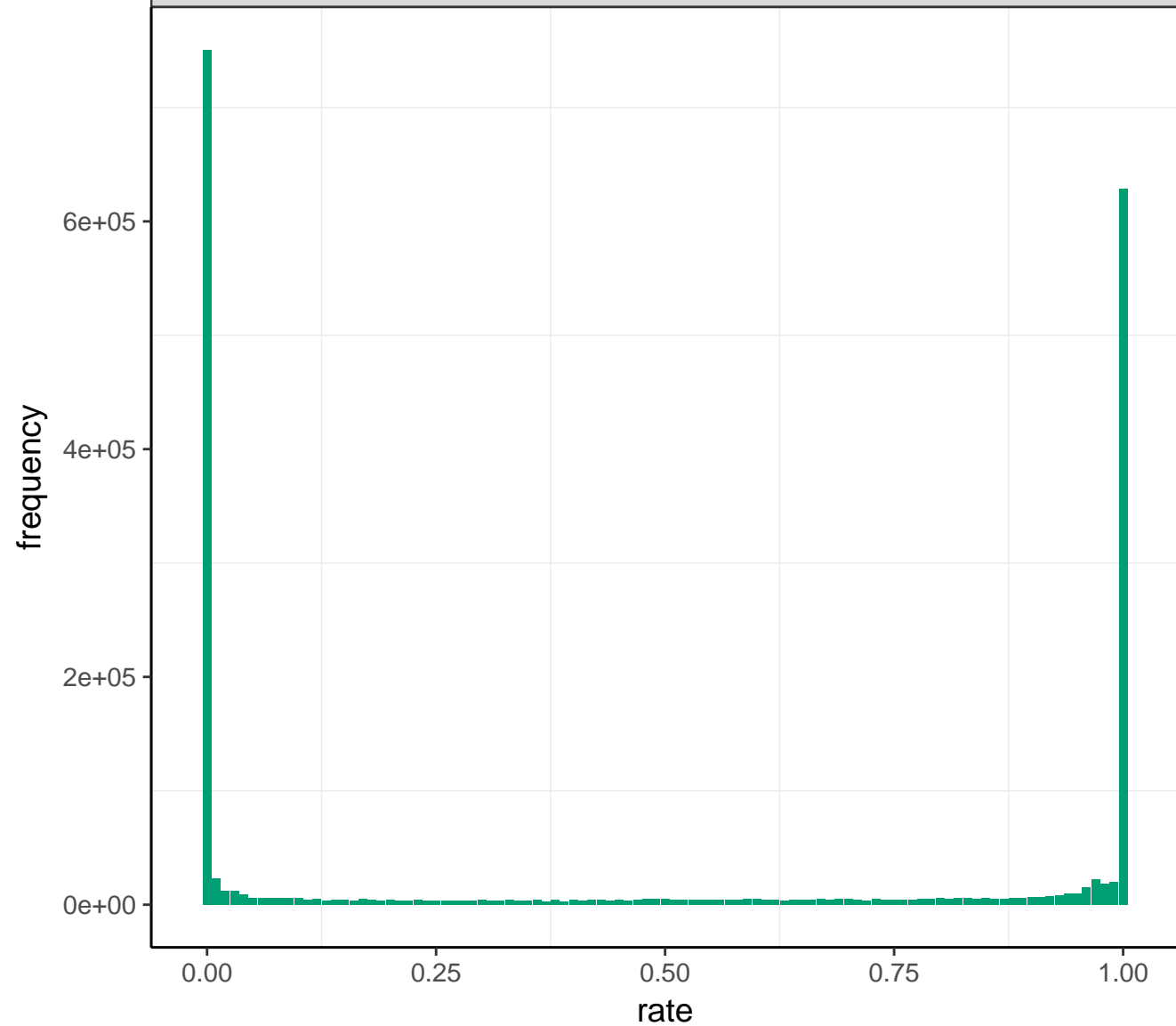
methylation rates at all positions

190626_I114_FCH7TVNBBXY_L3_5-S2

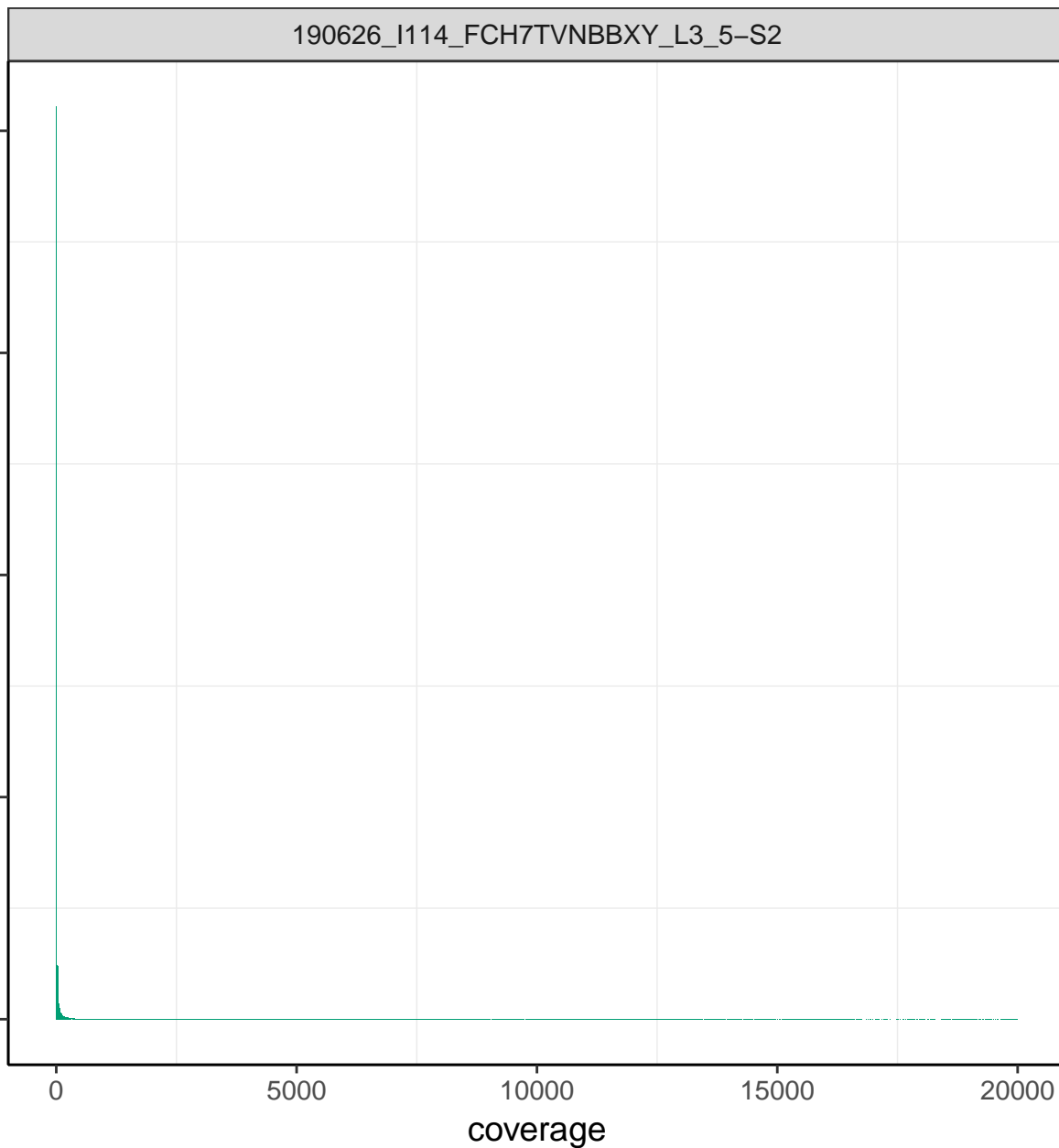


methylation rates at filtered positions
context:CG, min MDP:10, max MDP:500

190626_I114_FCH7TVNBBXY_L3_5-S2



coverages at all positions



coverages at all positions, in 1–1744 (90%–quantile)

190626_I114_FCH7TVNBBXY_L3_5–S2

frequency

4e+06

3e+06

2e+06

1e+06

0e+00

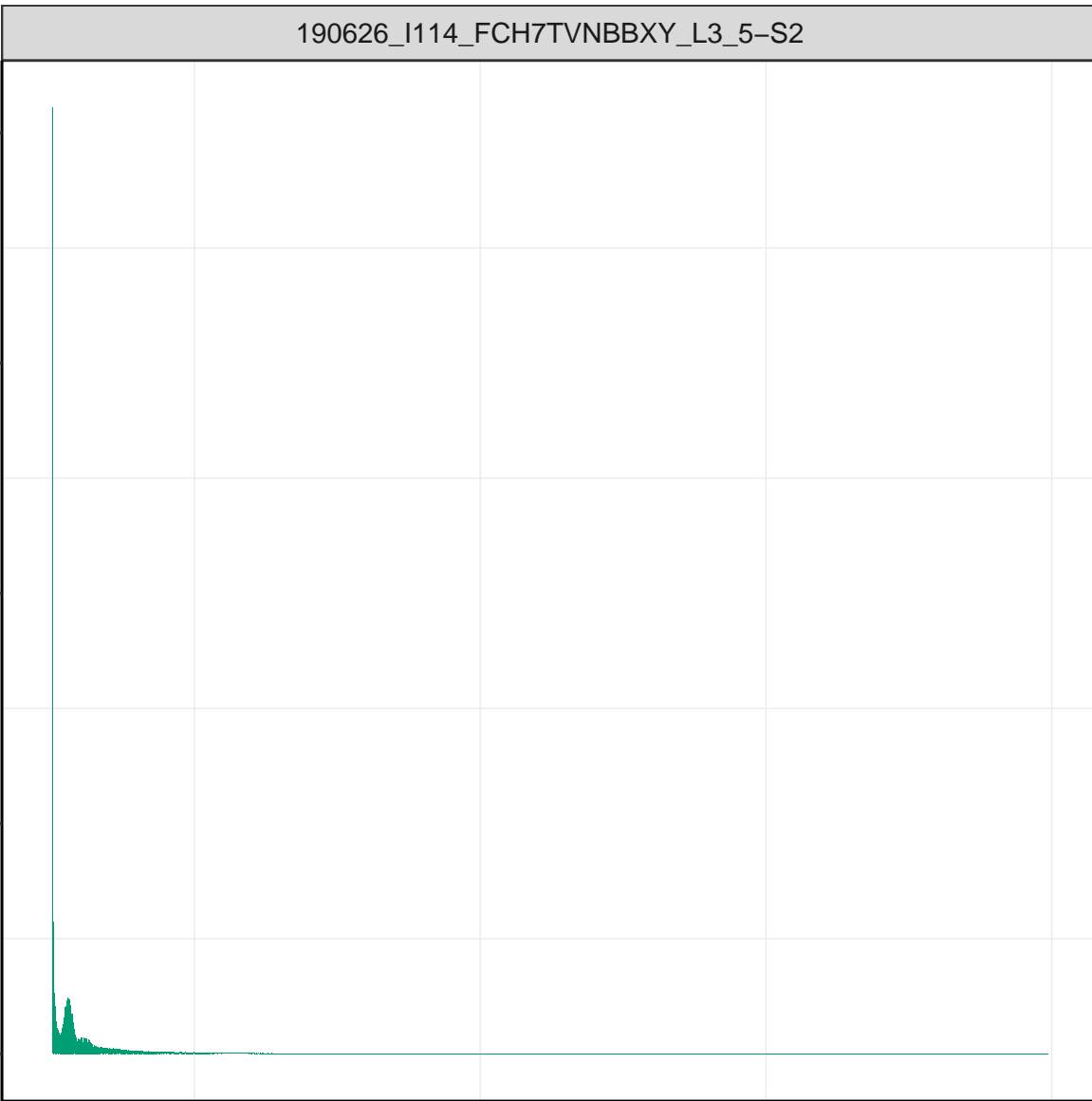
0

500

1000

1500

coverage



coverages at filtered positions
context:CG, min MDP:10, max MDP:500

190626_I114_FCH7TVNBBXY_L3_5-S2

