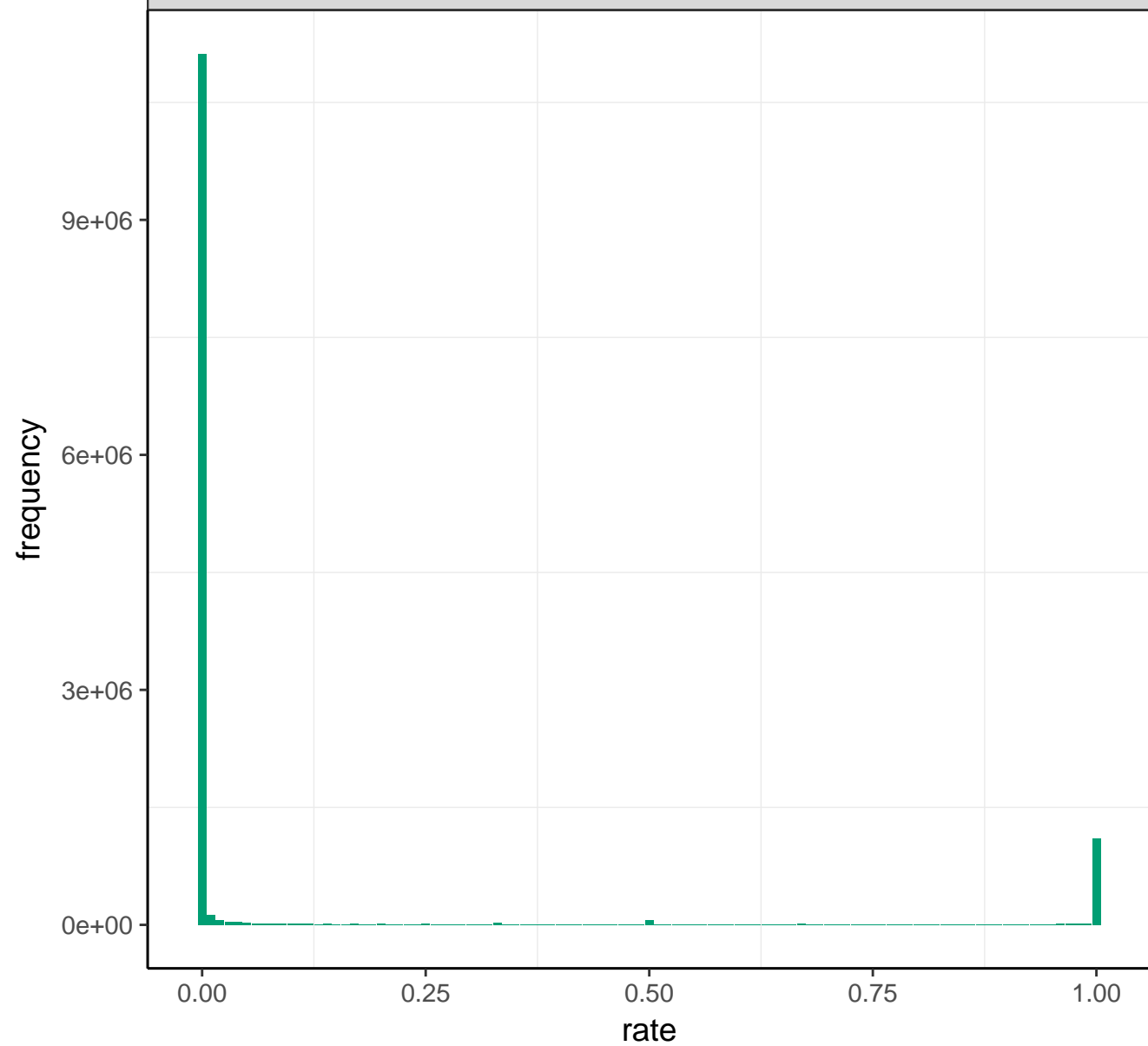
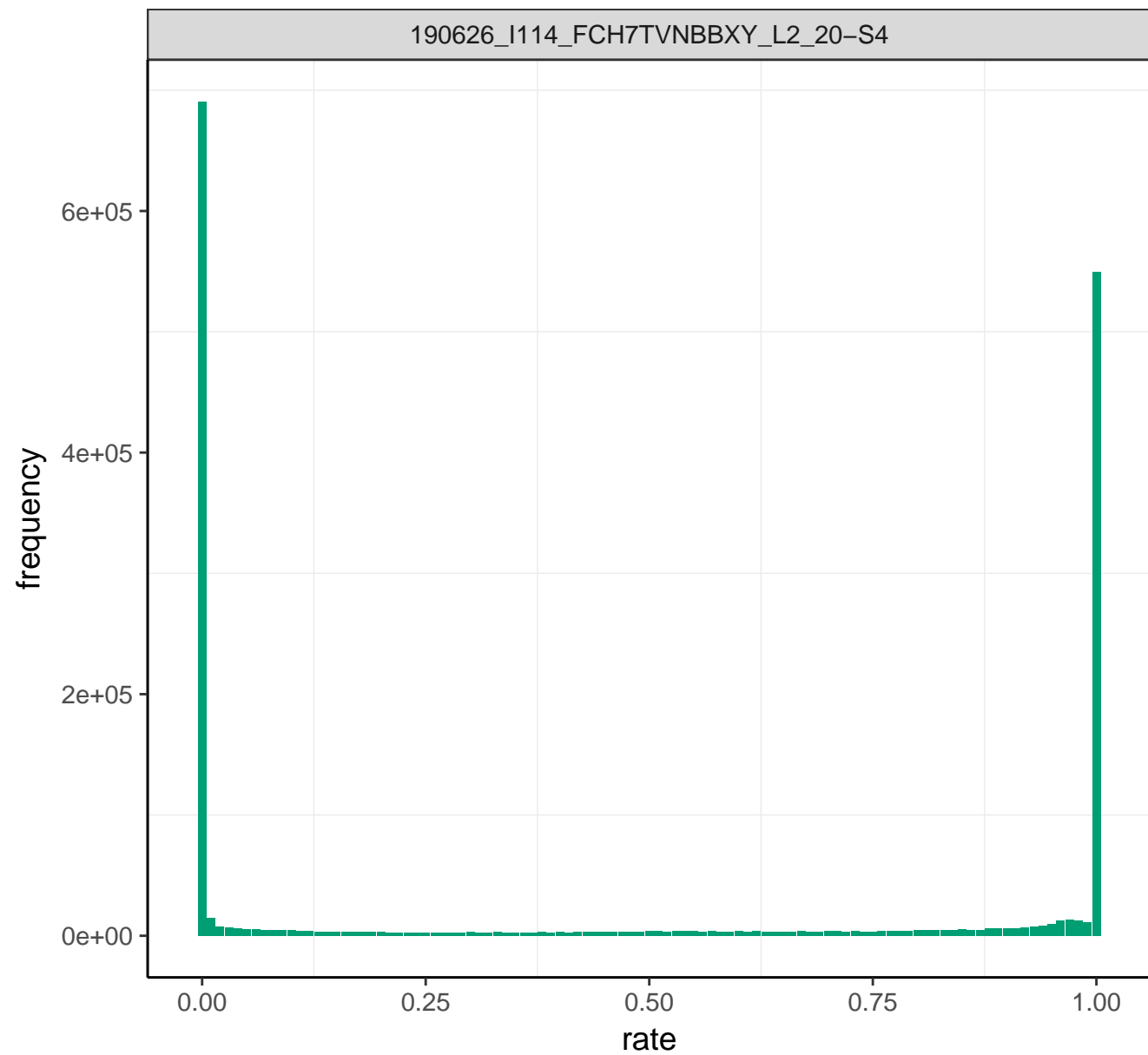


# methylation rates at all positions

190626\_I114\_FCH7TVNBBXY\_L2\_20-S4



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



# coverages at all positions

190626\_I114\_FCH7TVNBBXY\_L2\_20-S4

frequency

2e+06

1e+06

0e+00

0

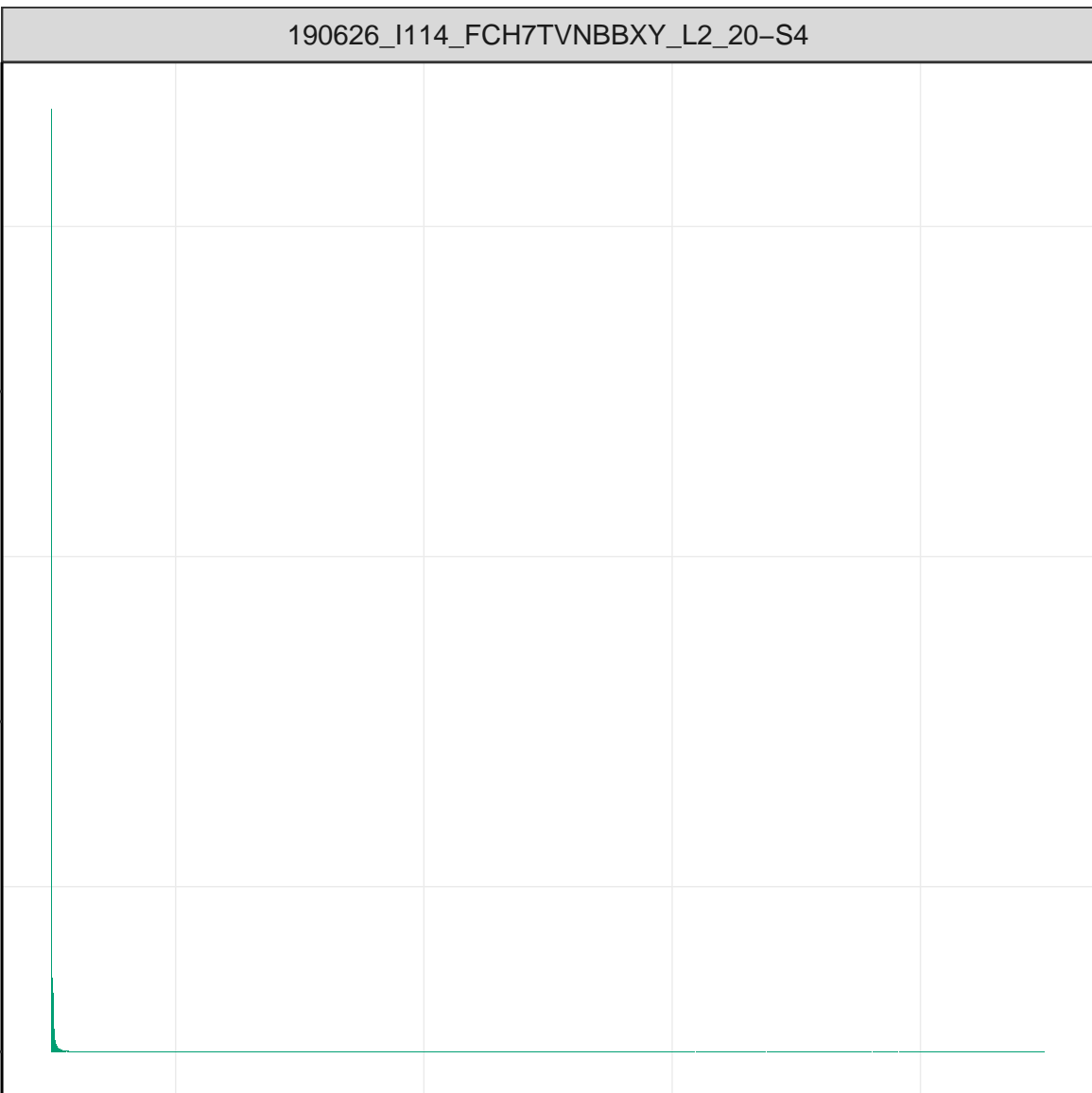
5000

10000

15000

20000

coverage



coverages at all positions, in 1–921 (90%–quantile)

190626\_I114\_FCH7TVNBBXY\_L2\_20–S4

frequency

2e+06

1e+06

0e+00

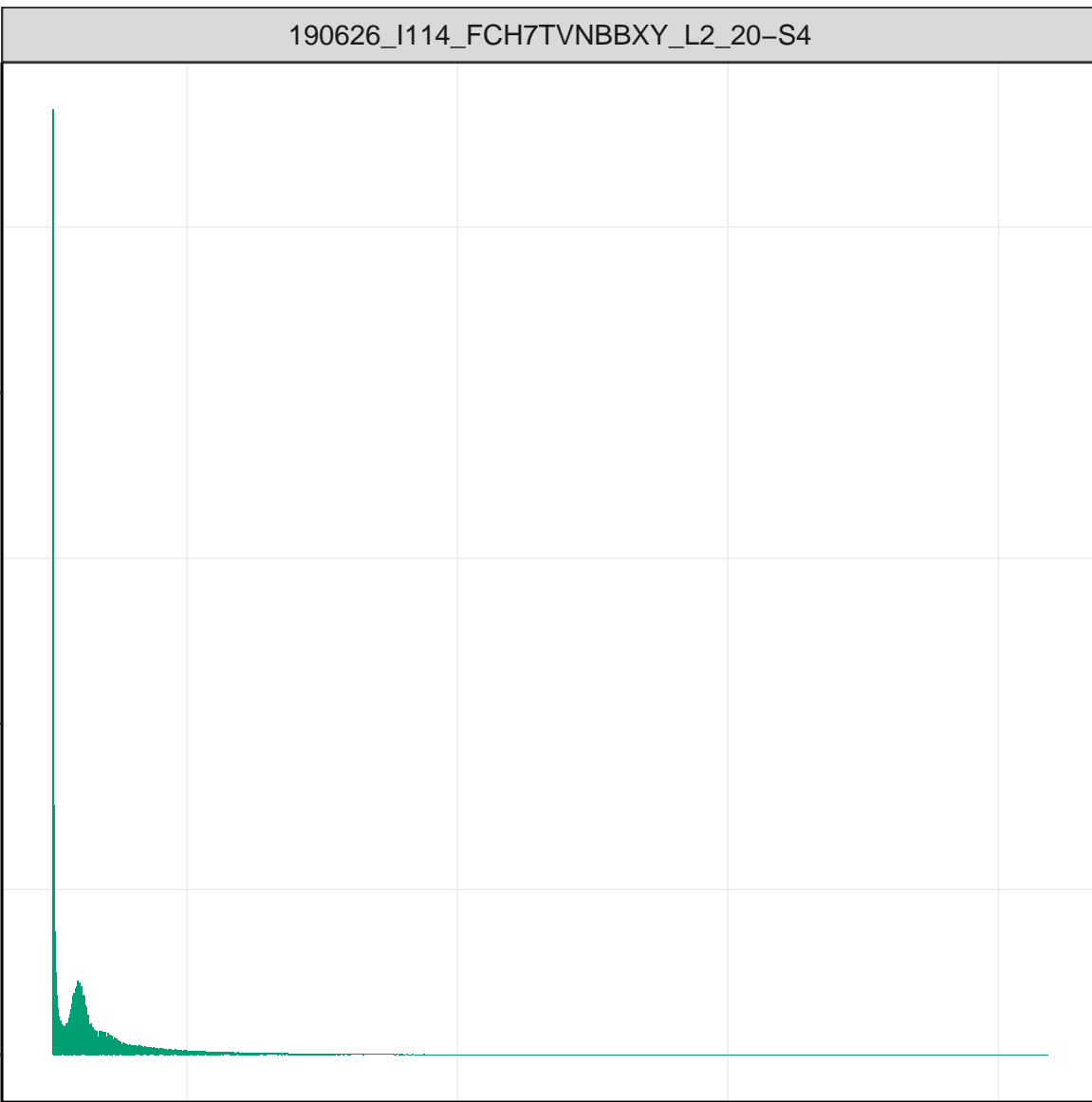
0

250

500

750

coverage



coverages at filtered positions

context:CG, min MDP:10

190626\_I114\_FCH7TVNBBXY\_L2\_20-S4

