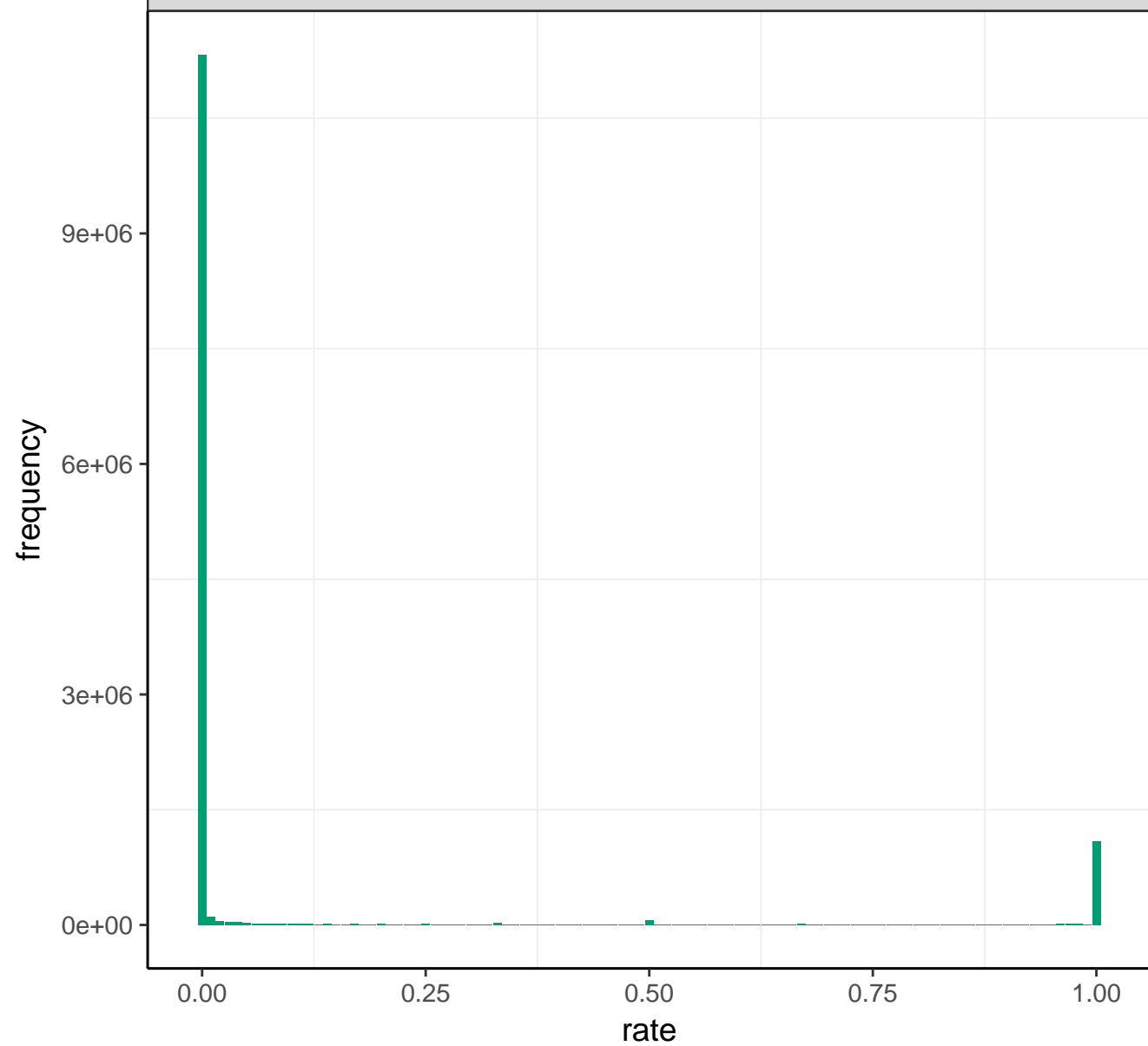
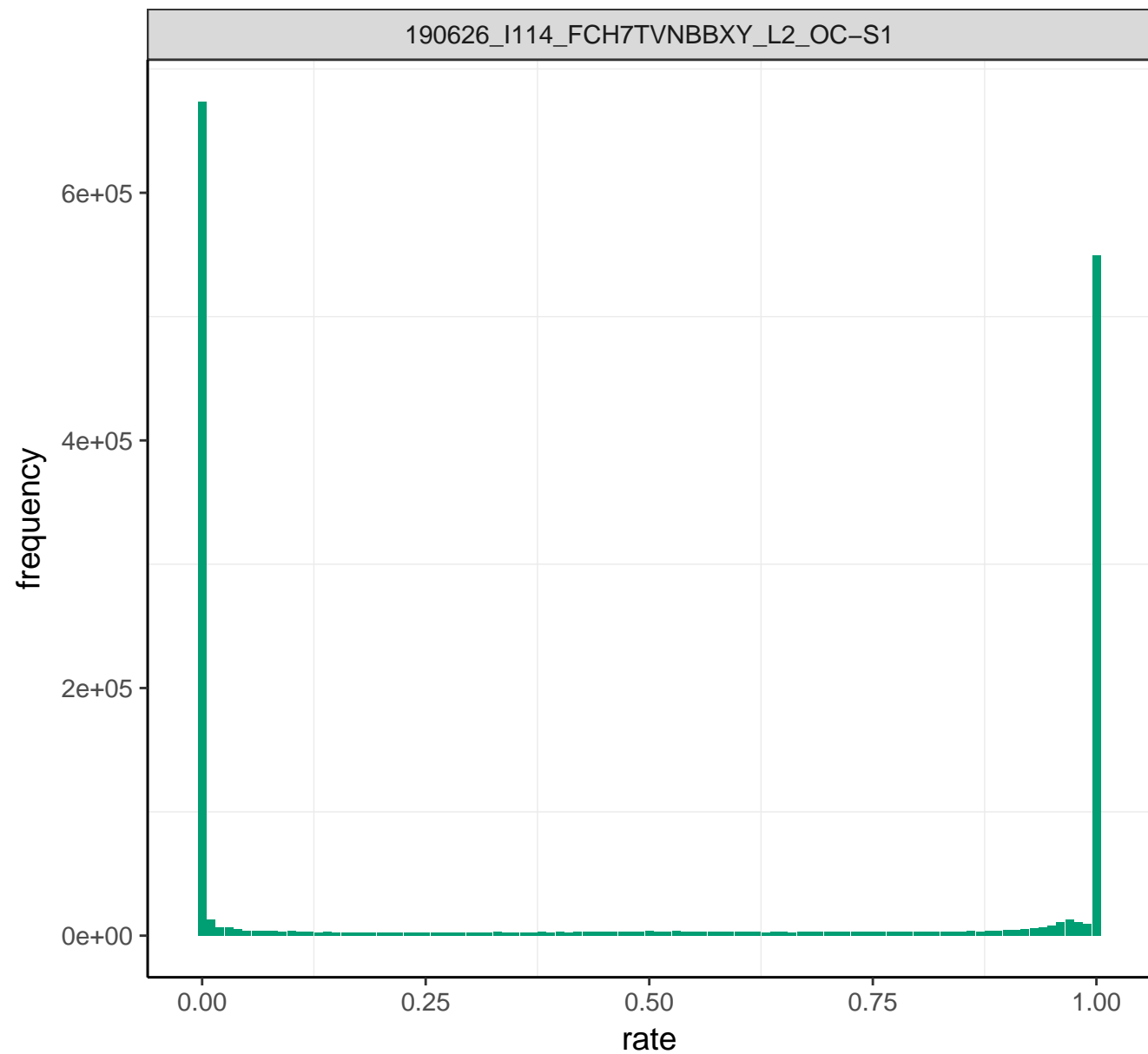


# methylation rates at all positions

190626\_I114\_FCH7TVNBBXY\_L2\_OC-S1



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



# coverages at all positions

190626\_I114\_FCH7TVNBBXY\_L2\_OC-S1

frequency

2e+06

1e+06

0e+00

0

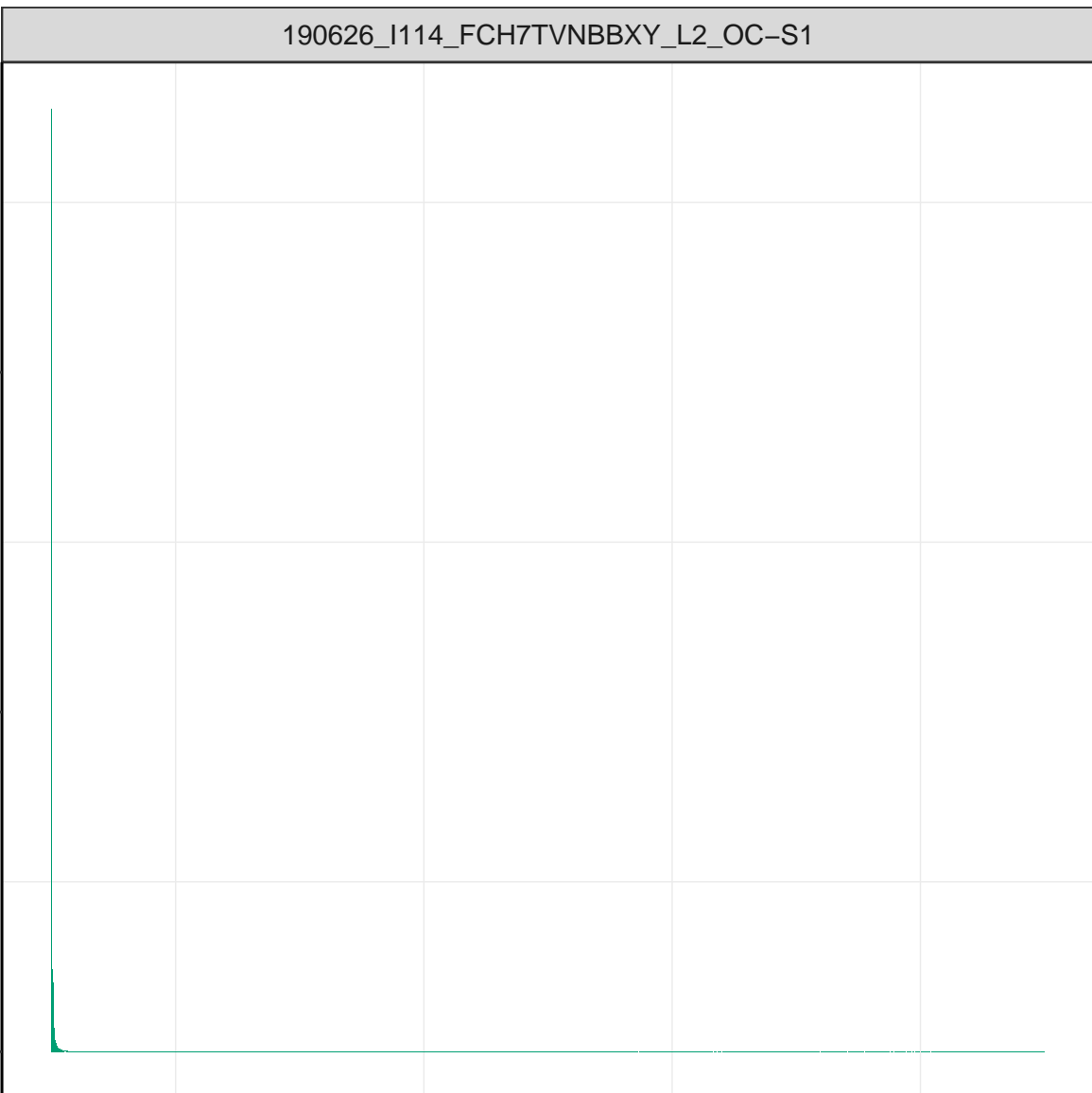
5000

10000

15000

20000

coverage



coverages at all positions, in 1–819 (90%–quantile)

190626\_I114\_FCH7TVNBBXY\_L2\_OC-S1

frequency

2e+06

1e+06

0e+00

0

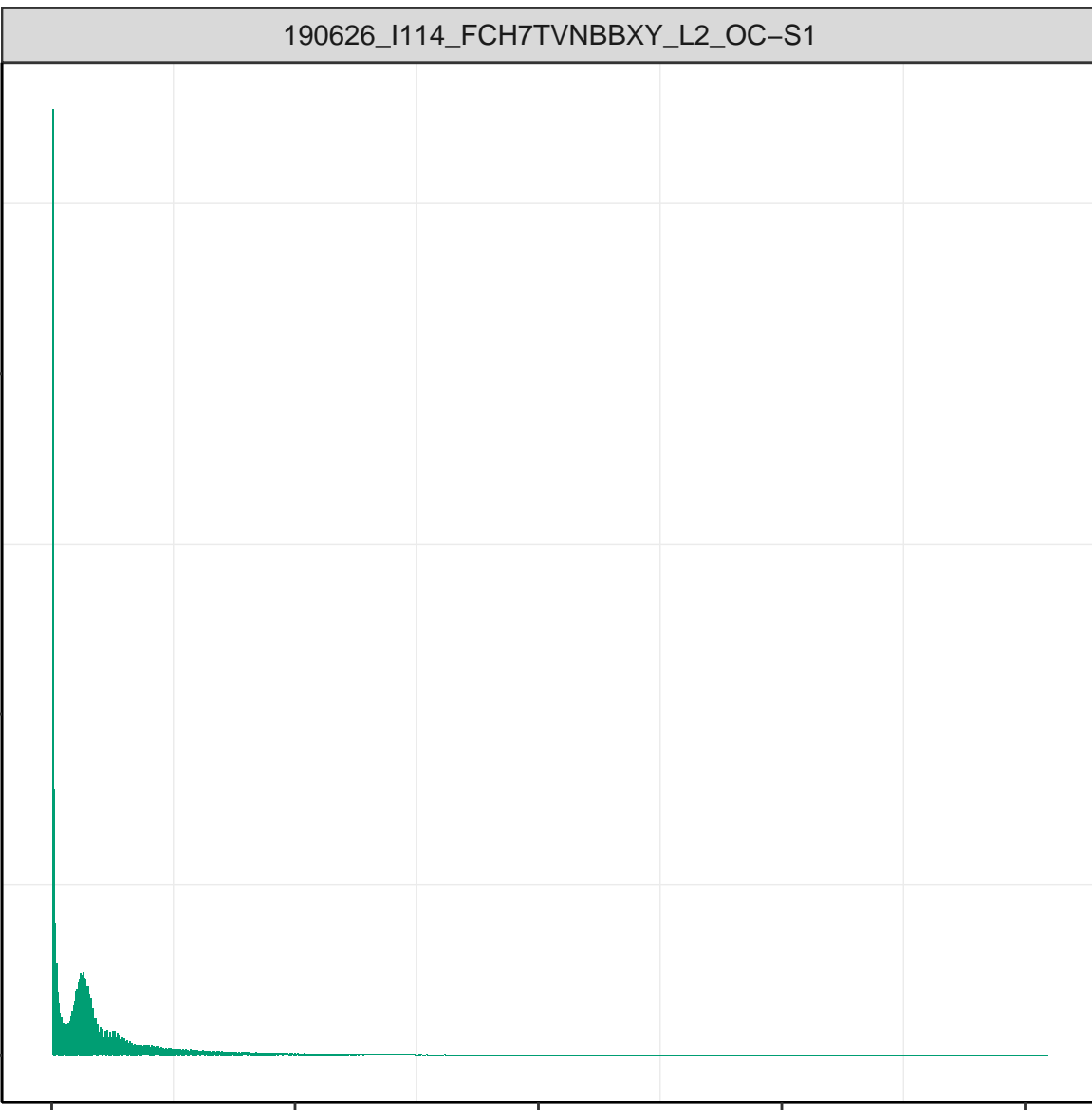
200

400

600

800

coverage



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

190626\_I114\_FCH7TVNBBXY\_L2\_OC-S1

