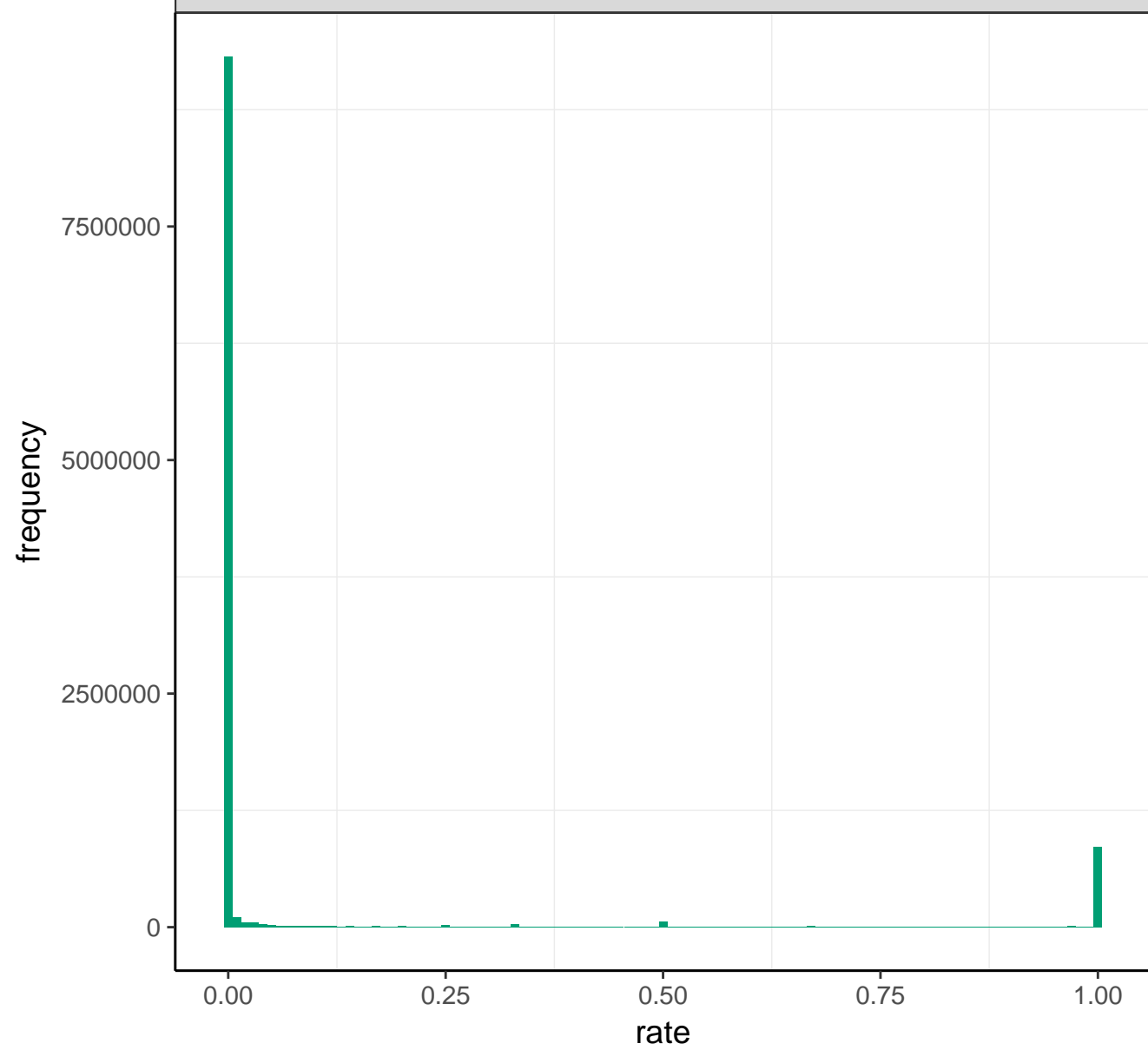
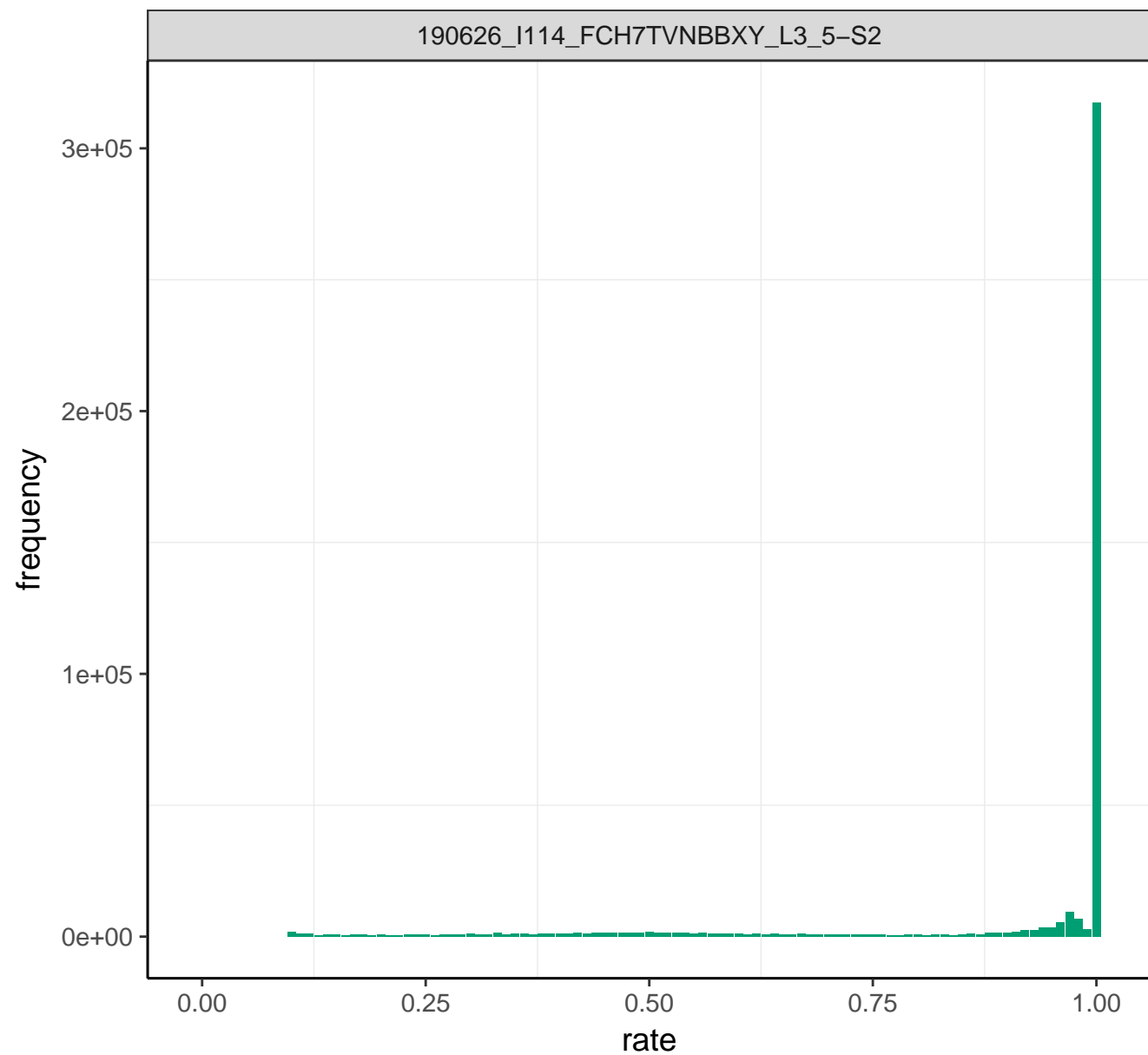


# methylation rates at all positions

190626\_I114\_FCH7TVNBBXY\_L3\_5-S2



methylation rates at filtered positions  
context:CG, min MDP:10, max MDP:100, min MR:0.1



# coverages at all positions

190626\_I114\_FCH7TVNBBXY\_L3\_5-S2

frequency

2e+06

1e+06

0e+00

0

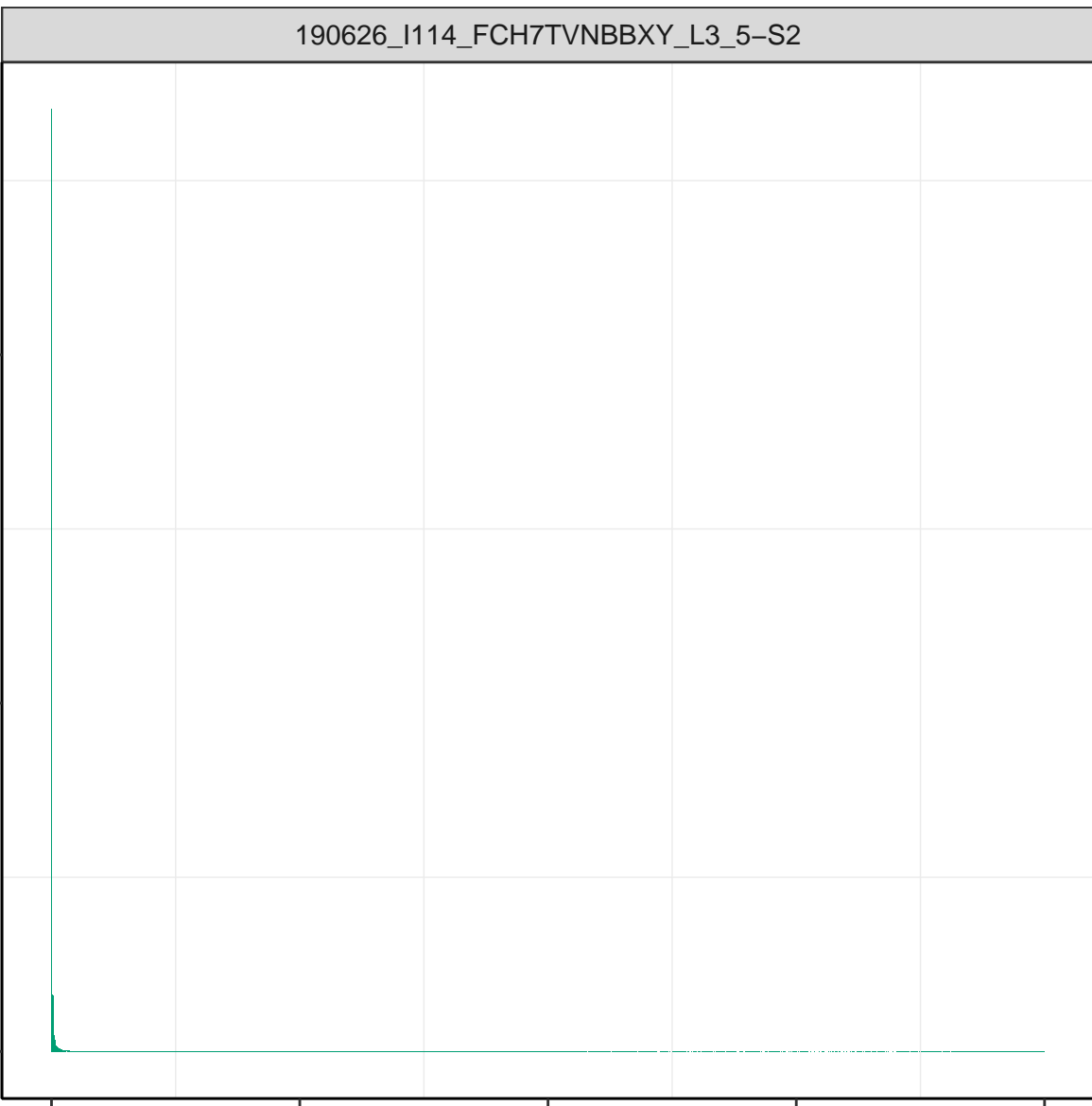
5000

10000

15000

20000

coverage



coverages at all positions, in 1–872 (90%–quantile)

190626\_I114\_FCH7TVNBBXY\_L3\_5–S2

frequency

2e+06

1e+06

0e+00

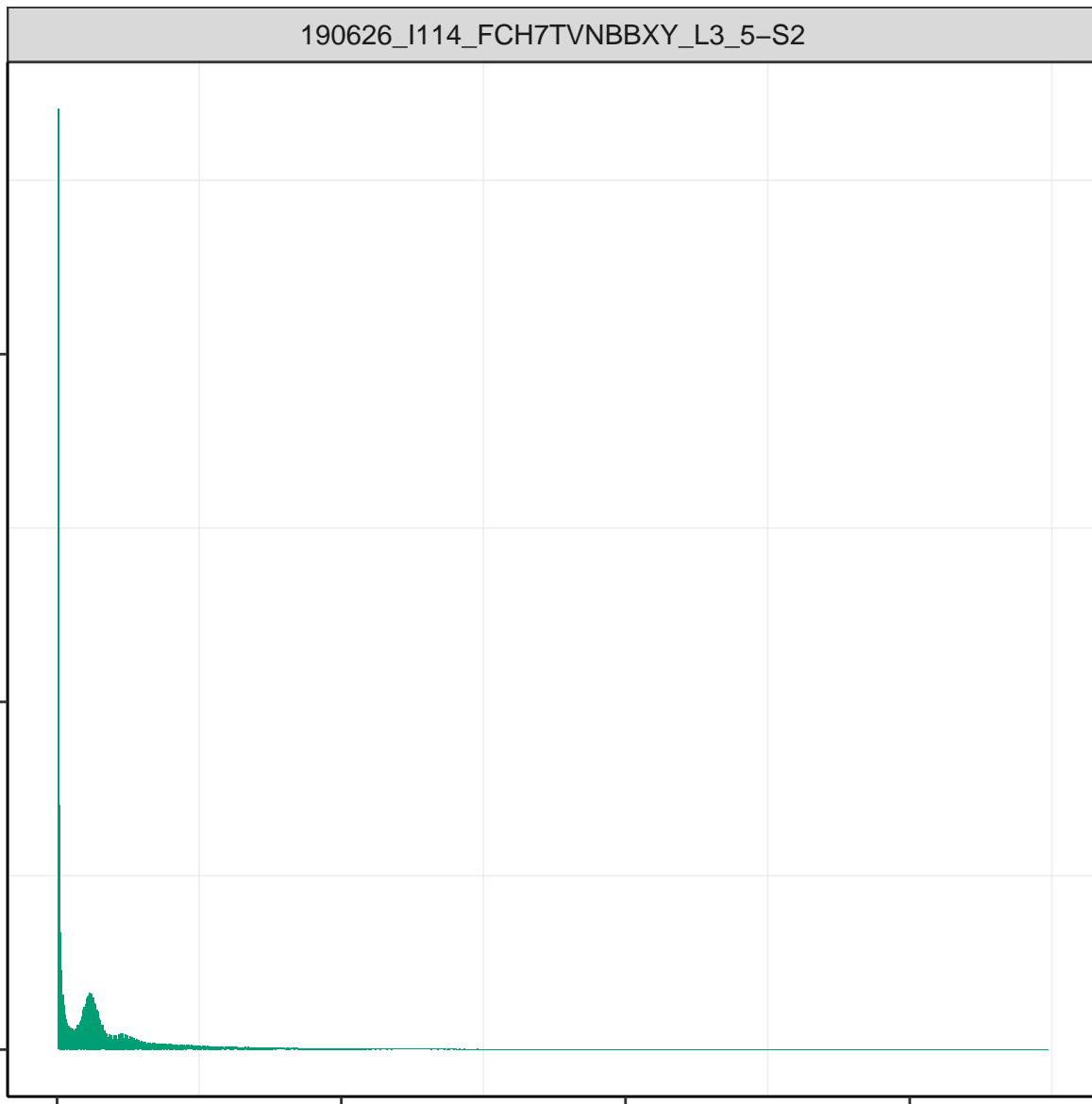
0

250

500

750

coverage



coverages at filtered positions

context:CG, min MDP:10, max MDP:100, min MR:0.1

190626\_I114\_FCH7TVNBBXY\_L3\_5-S2

