

GTseq.prod.all

Mean Coverage at Target SNP

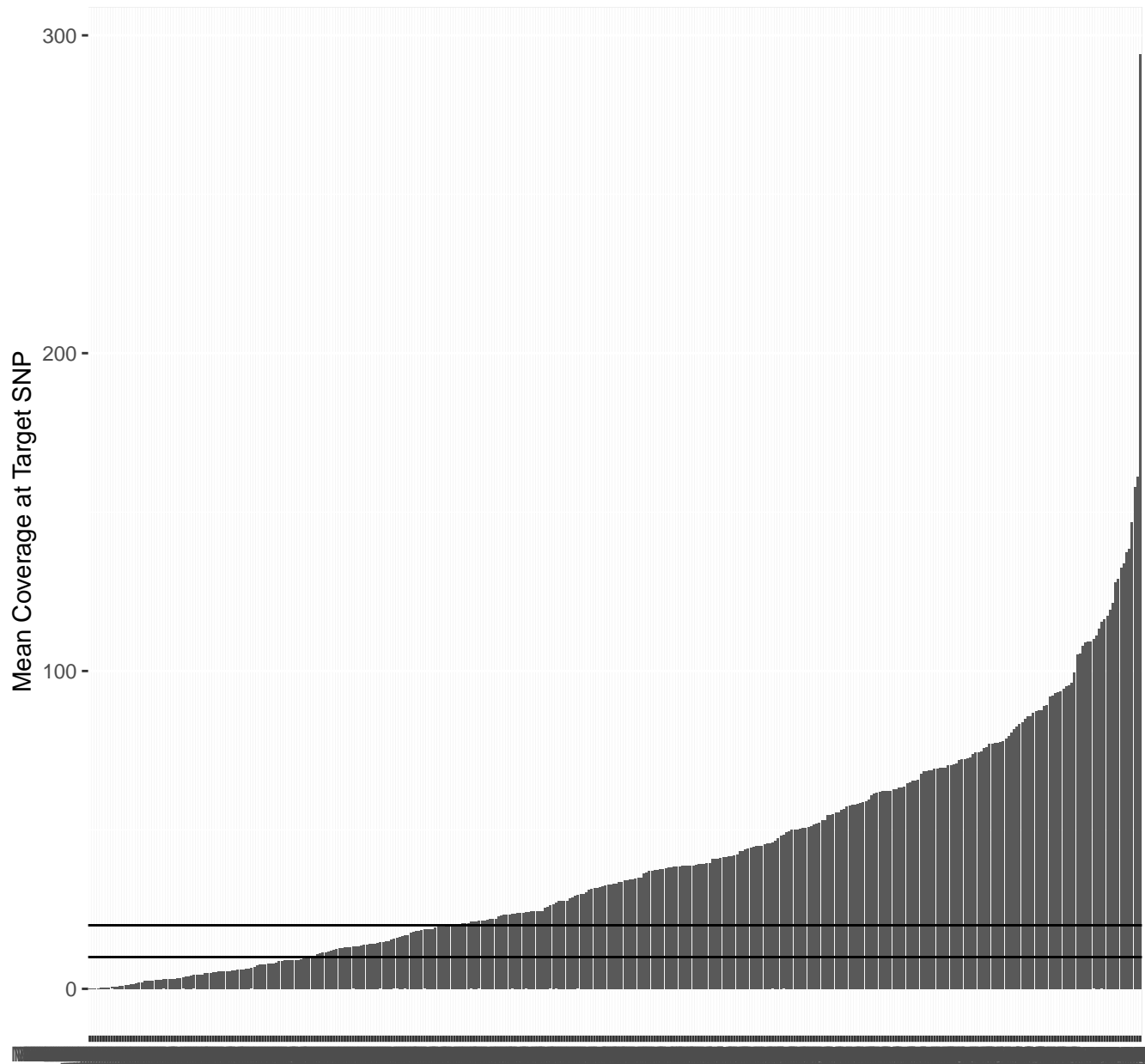
300 -

200 -

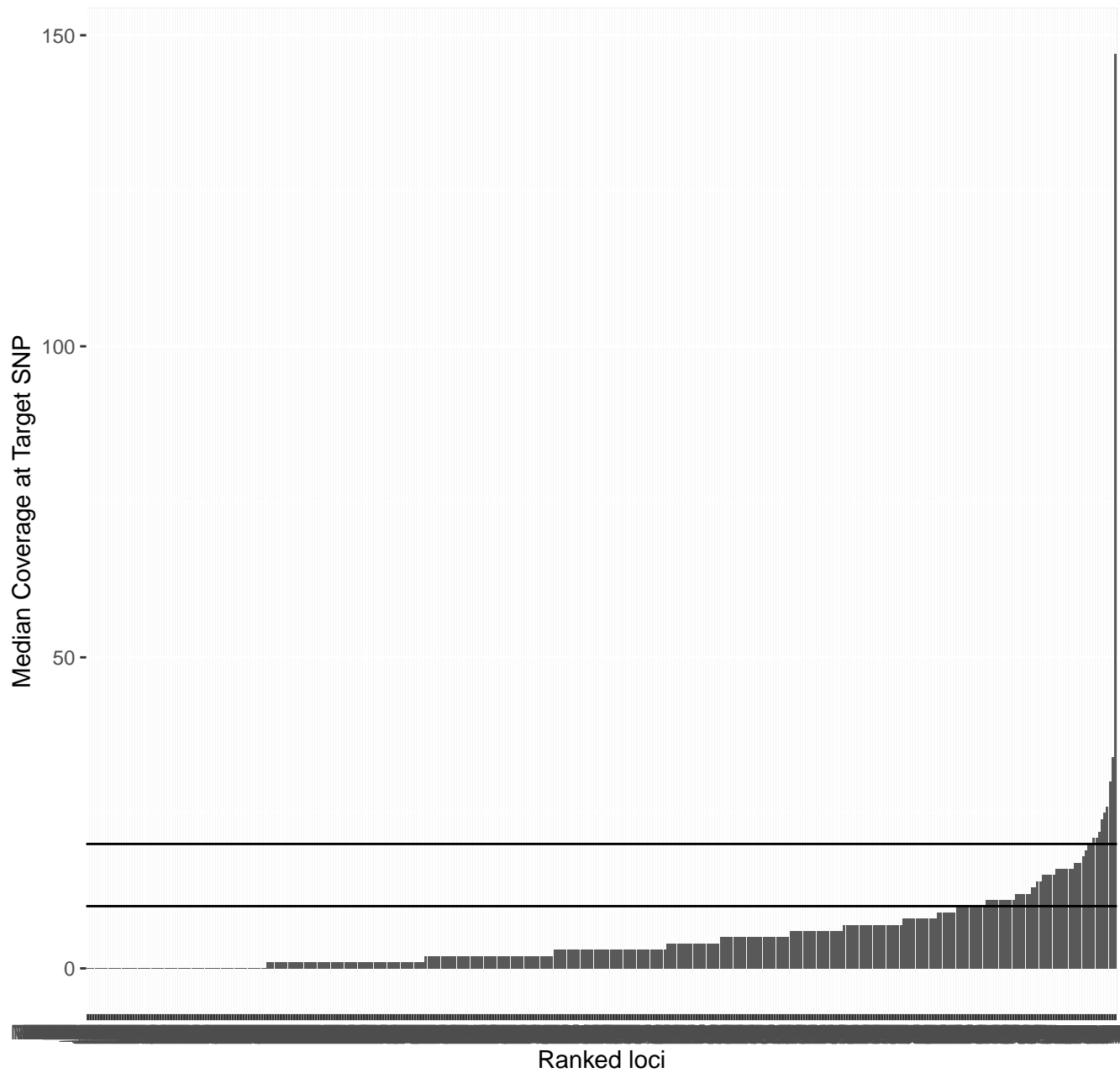
100 -

0

Ranked loci



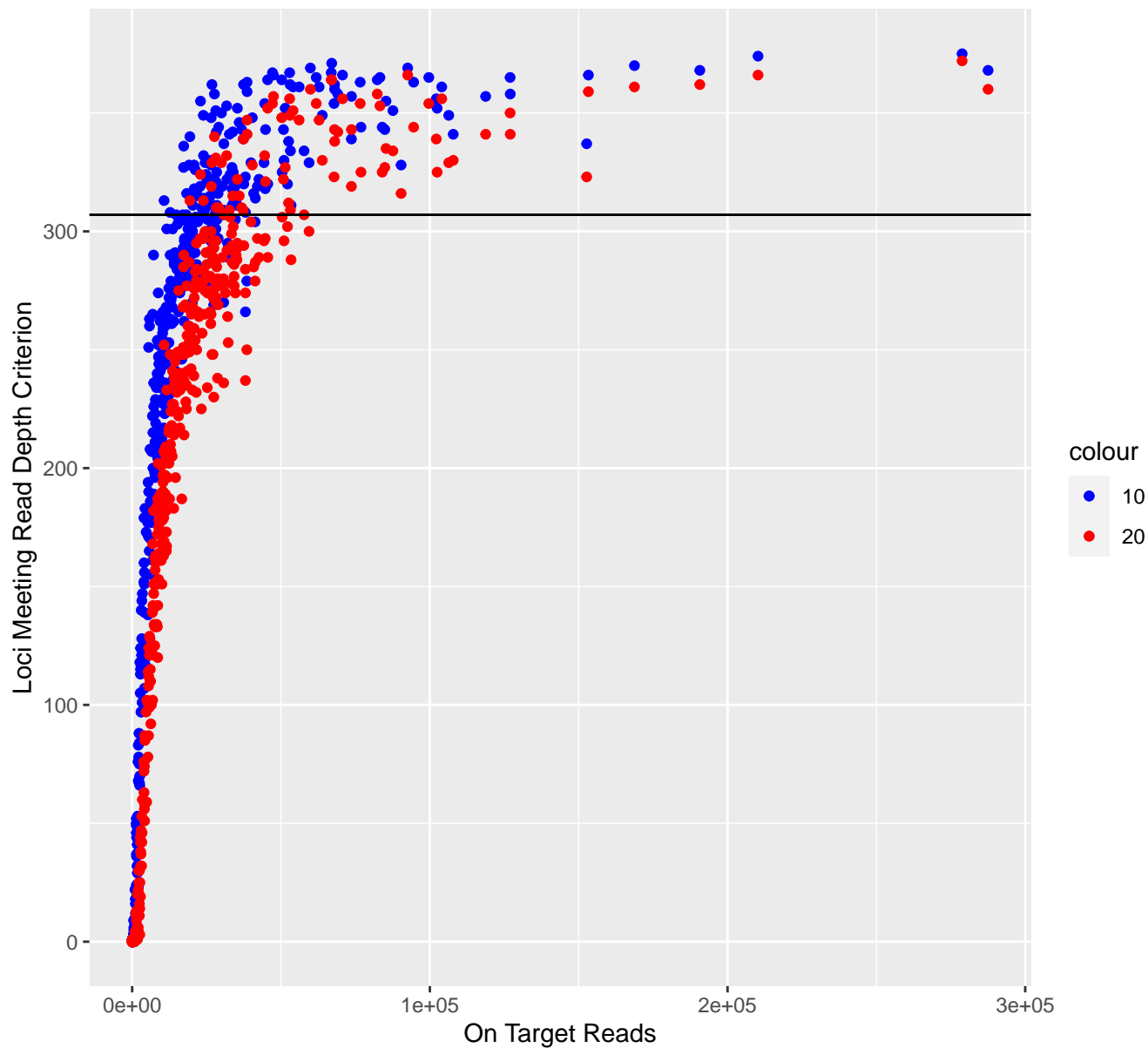
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Num individuals with read depth ≥ 10 at 307 loci: 152

Num individuals with read depth ≥ 20 at 307 loci: 77



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