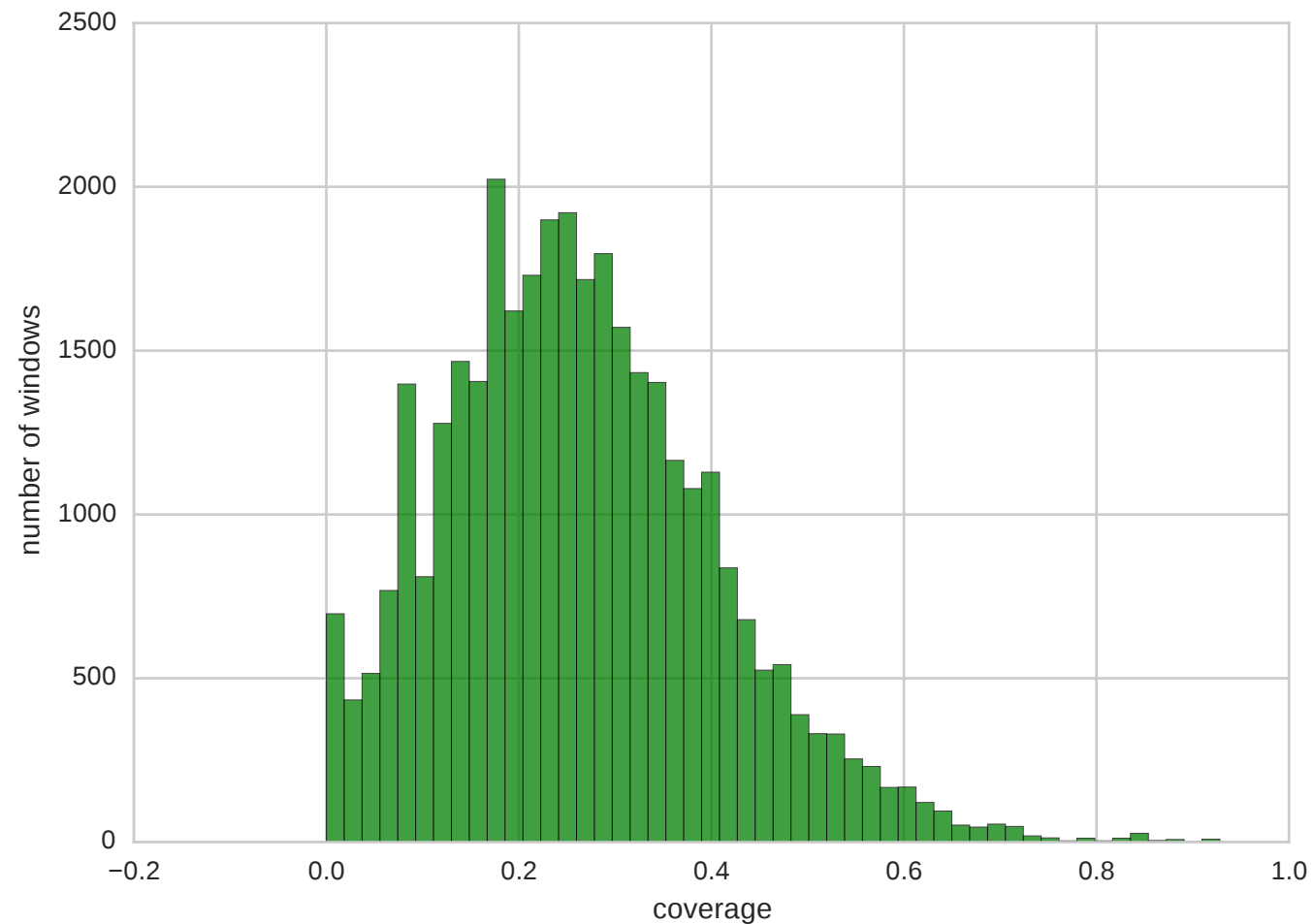
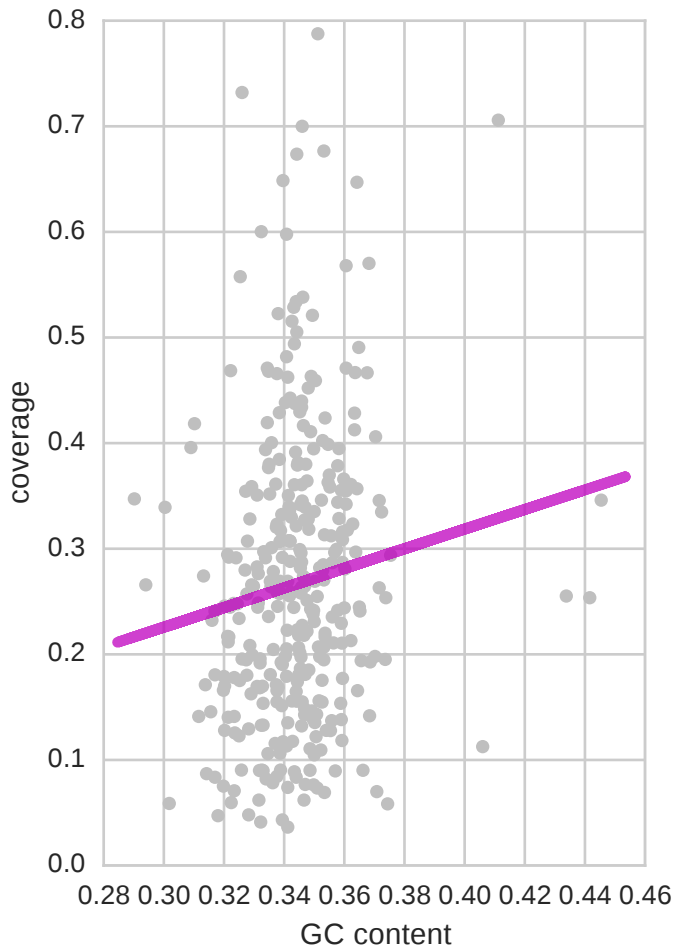


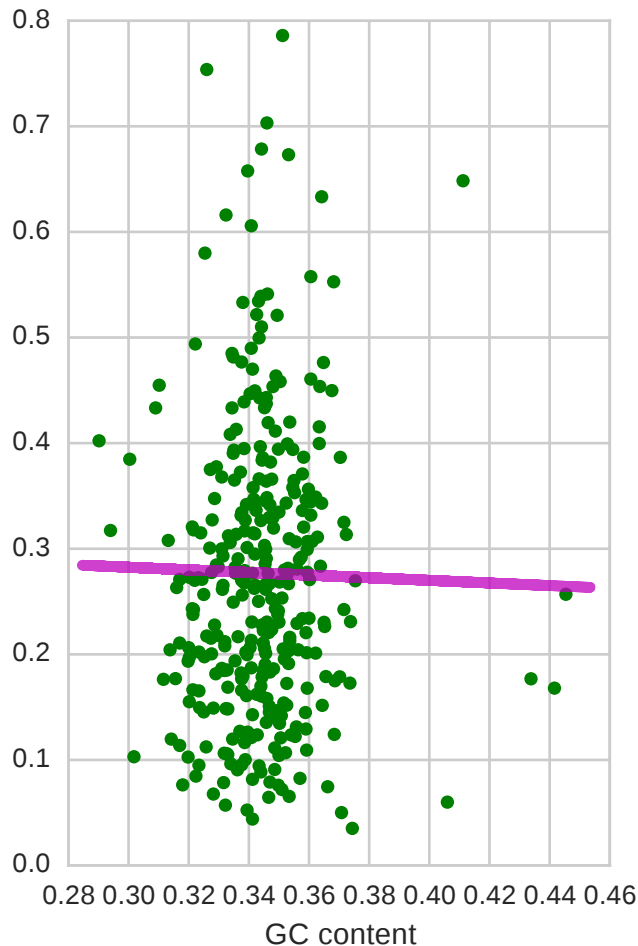
genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-64
iRep: n/a r^2 : 0.97 avg. cov: 0.26 % windows: 97.07



genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-64
iRep: n/a raw iRep: 5.34 GC r²: 0.02 GC bias: 0.02 avg. cov: 0.26 % windows: 97.07

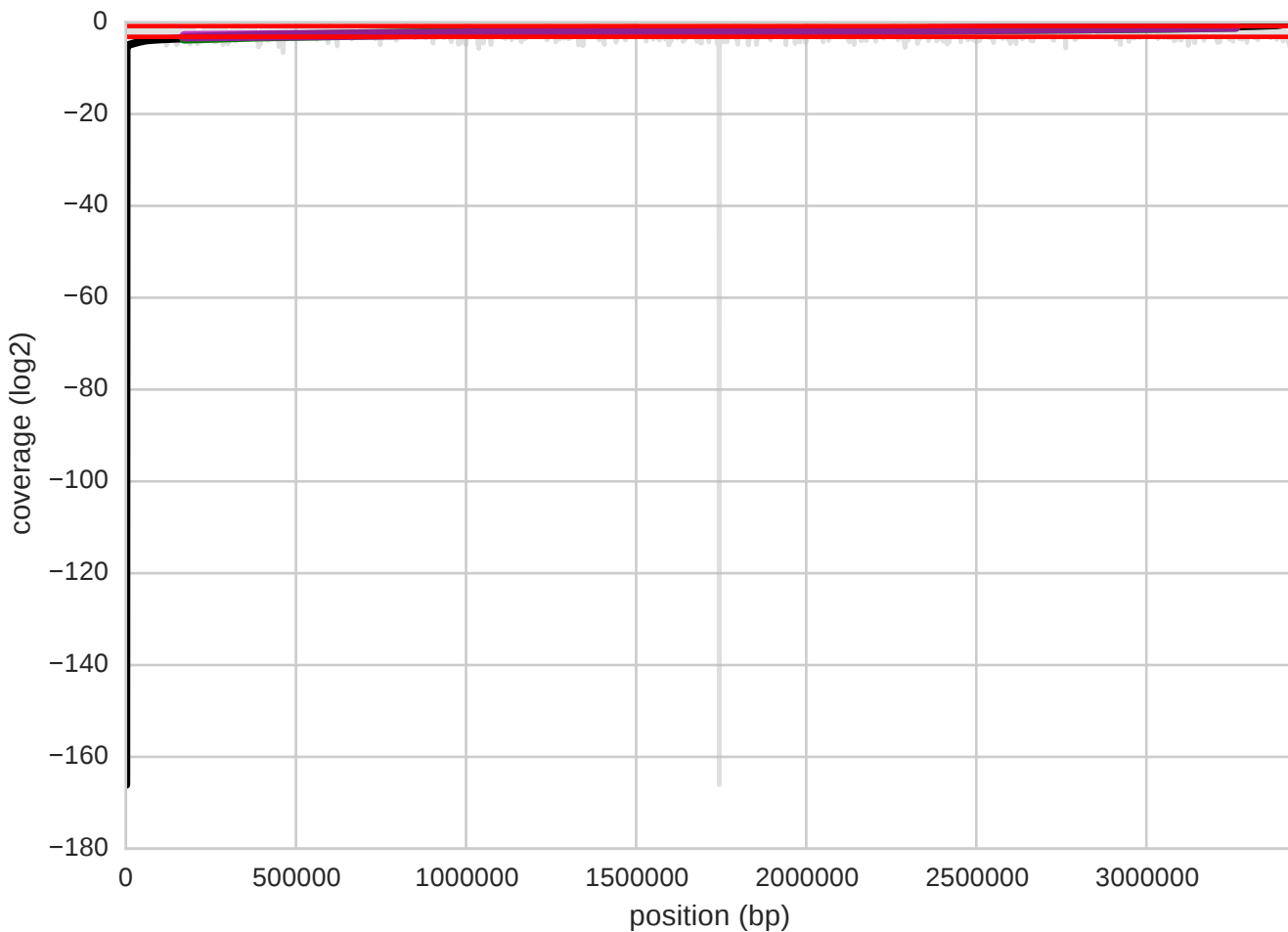


● filtered coverage
— least squares fit to GC filtered data



● corrected
— least squares fit to corrected data

genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-64
iRep: n/a un-filtered iRep: 5.04 r^2 : 0.97 avg. cov: 0.26 % windows: 97.07



— filtered
— sorted
— trimmed
— least squares fit to trimmed
— Ori/Ter