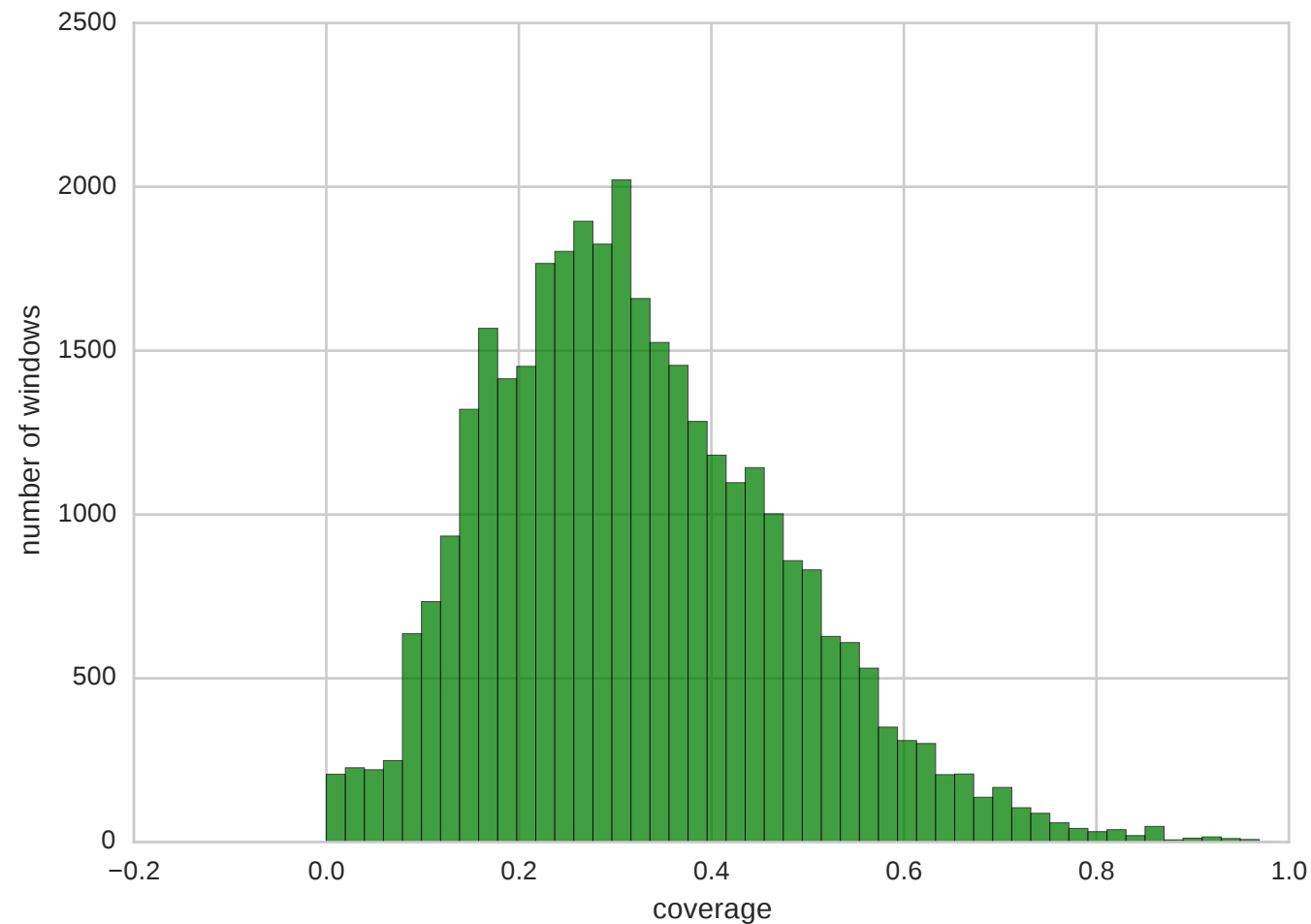
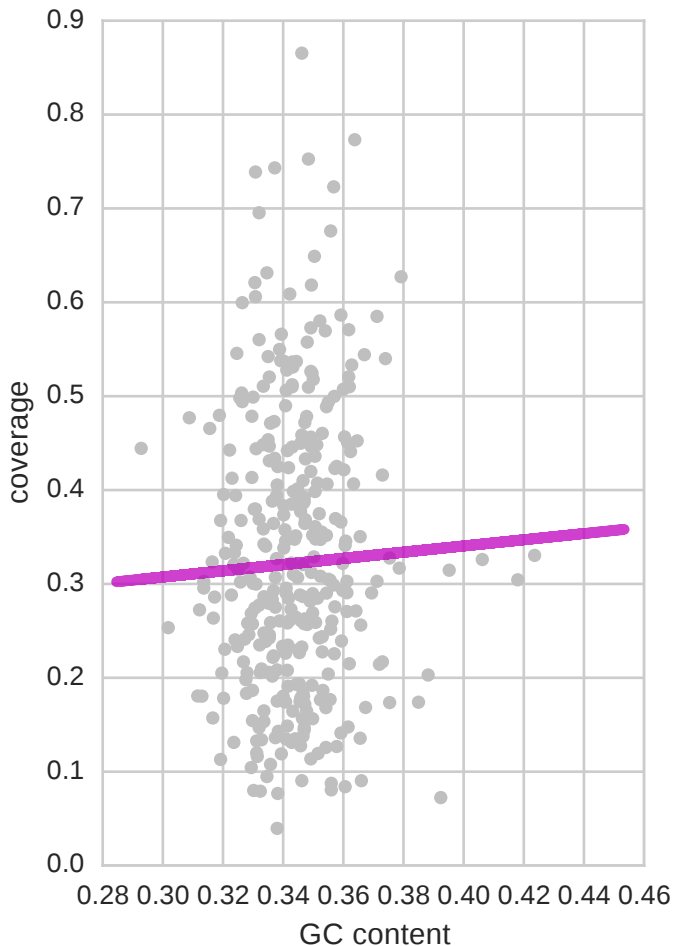


genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-65
iRep: n/a r^2 : 0.98 avg. cov: 0.32 % windows: 98.75

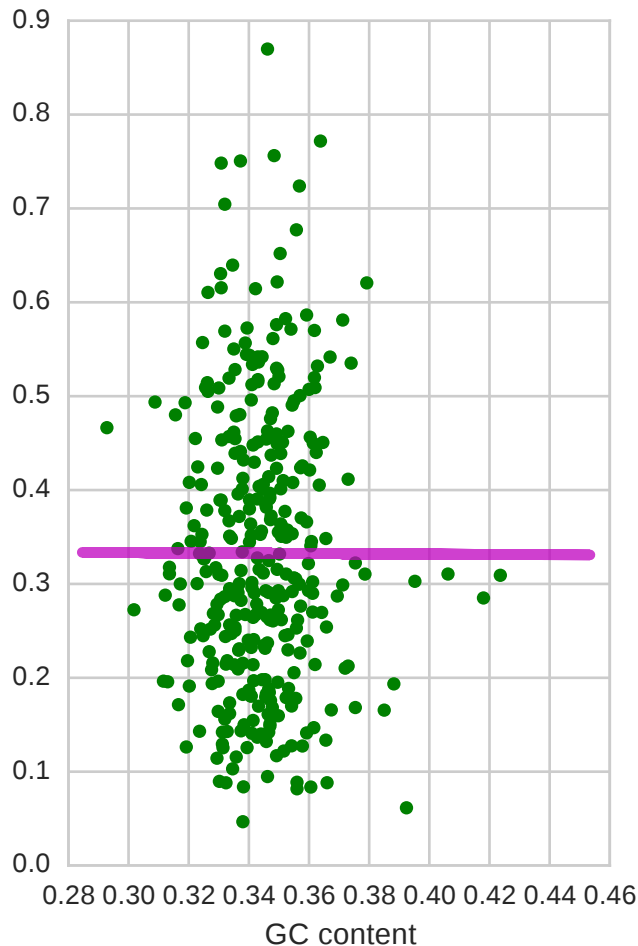


genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-65
iRep: n/a raw iRep: 4.52 GC r²: 0.00 GC bias: 0.00 avg. cov: 0.32 % windows: 98.75



● filtered coverage

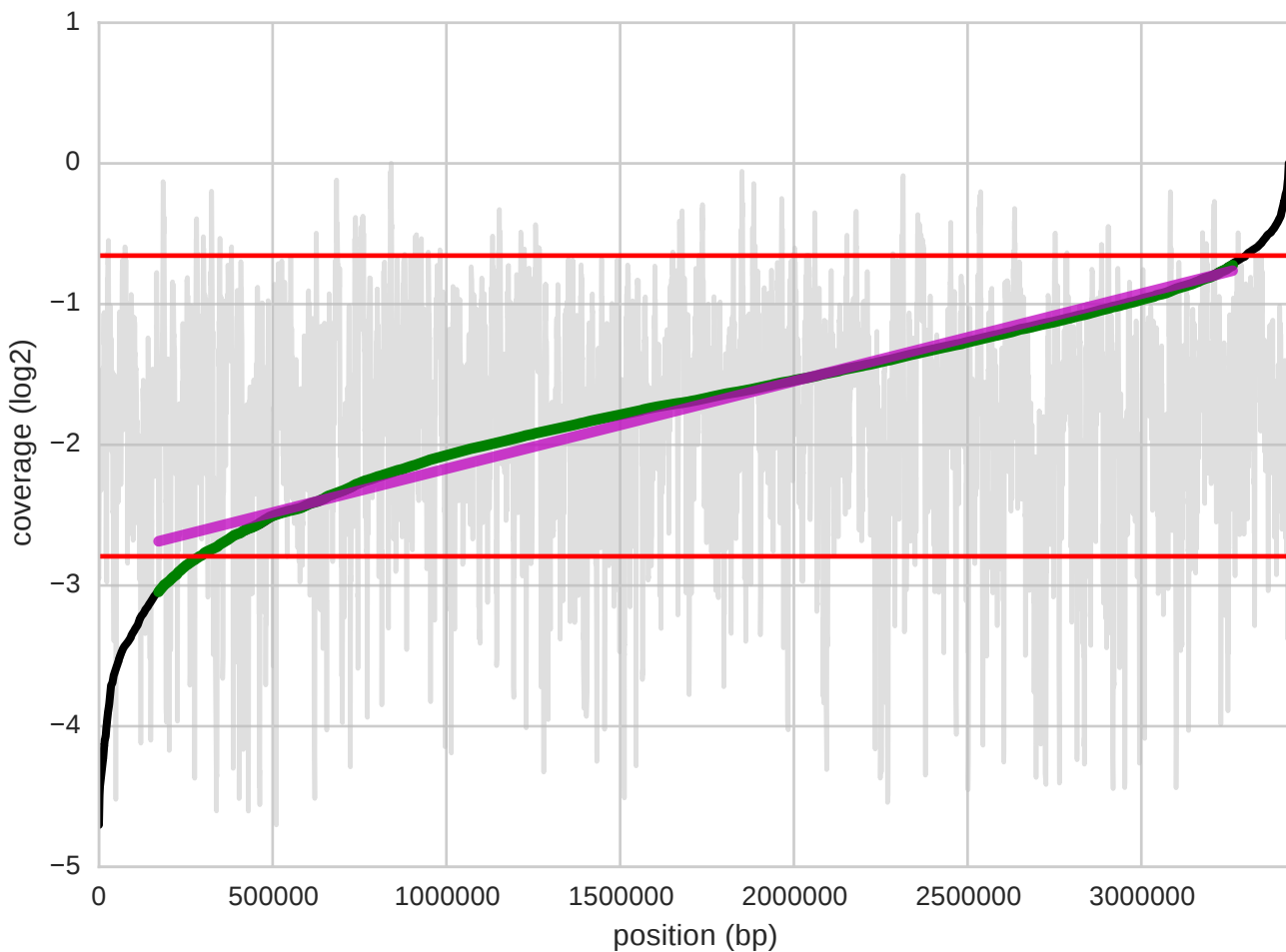
— least squares fit to GC filtered data



● corrected

— least squares fit to corrected data

genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-65
iRep: n/a un-filtered iRep: 4.40 r²: 0.98 avg. cov: 0.32 % windows: 98.75



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