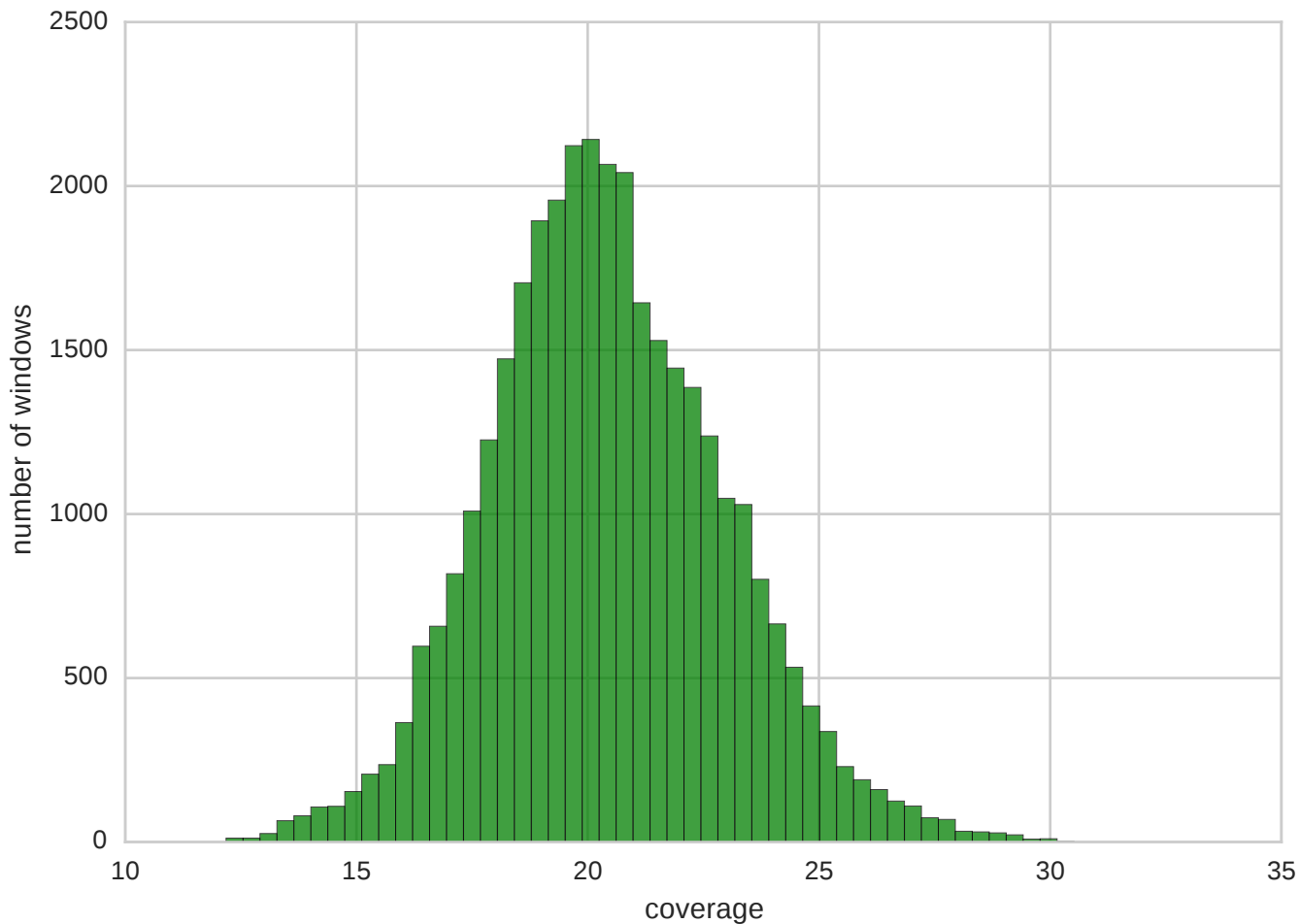
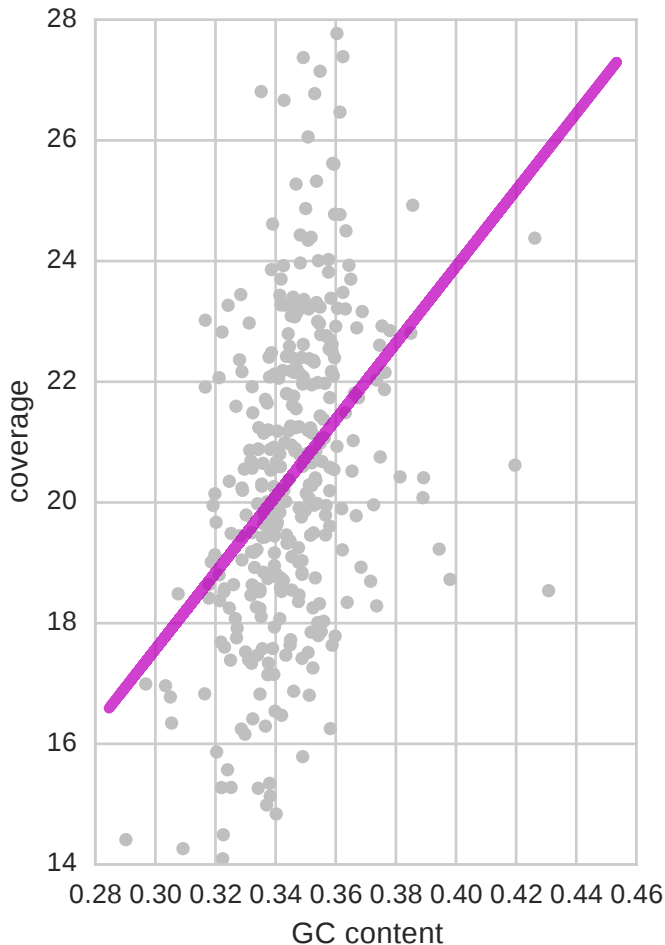


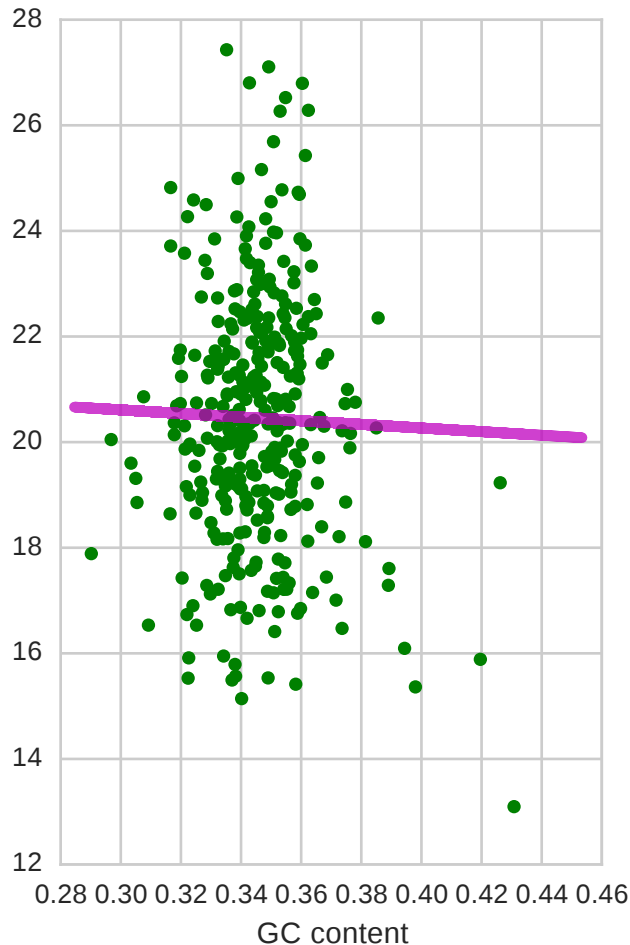
genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-17
iRep: 1.40 r²: 0.99 avg. cov: 20.42 % windows: 100.00



genome: ./bins/bacla_vizbin2-contigs sample: ./splitBam/bacla_vizbin2/DNA-Pieter-17
iRep: 1.40 raw iRep: 1.45 GC r²: 0.20 GC bias: 12.38 avg. cov: 20.42 % windows: 100.00



● 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-17
iRep: 1.40 un-filtered iRep: 1.40 r^2 : 0.99 avg. cov: 20.42 % windows: 100.00

