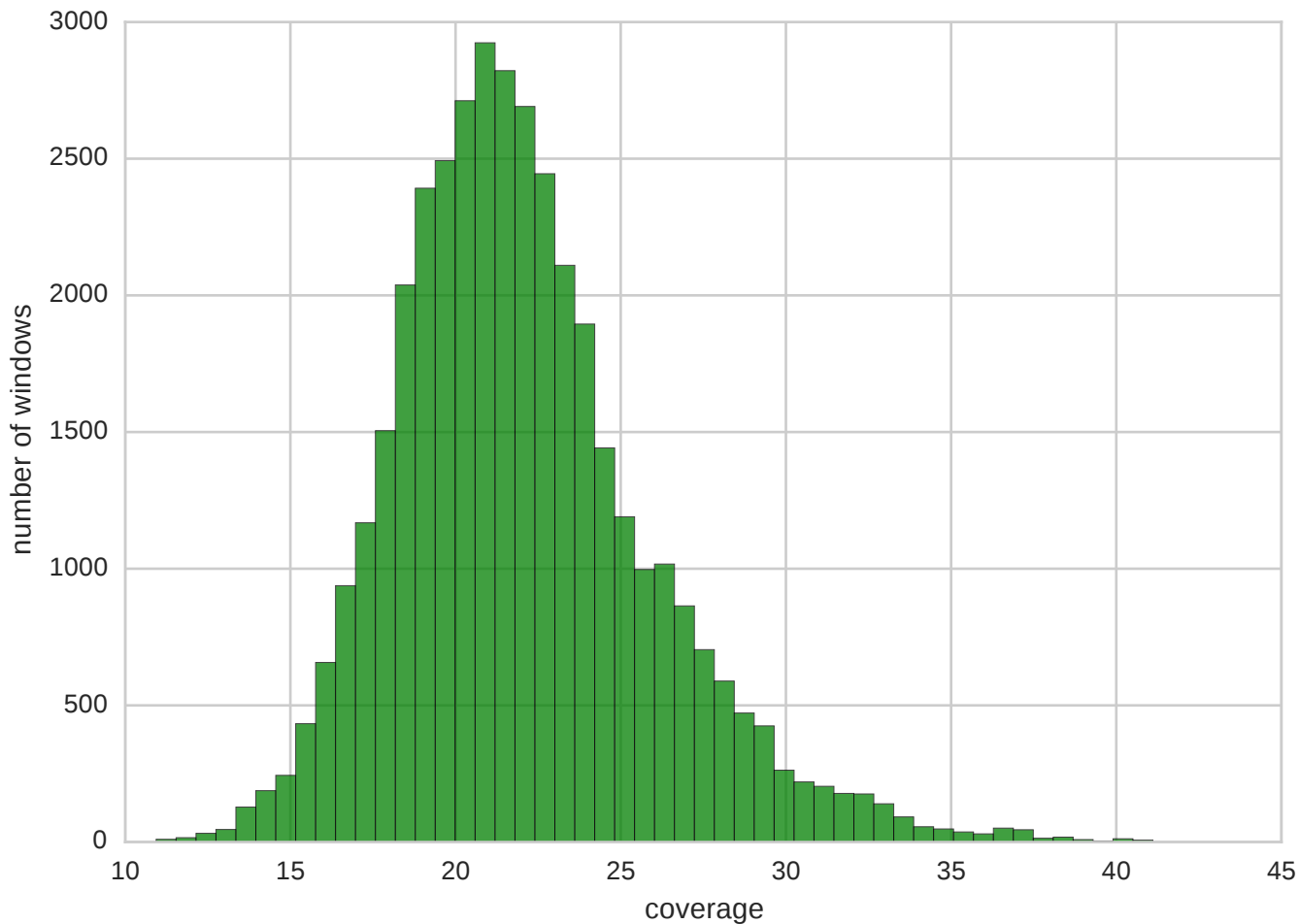
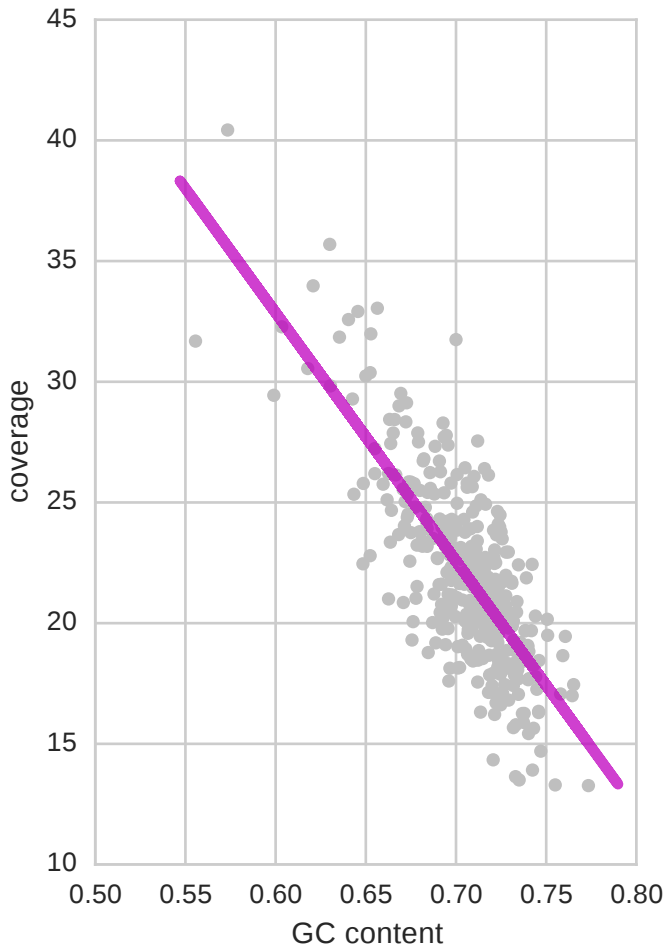


genome: ./bins/Betla_bin-contigs sample: ./splitBam/Betla_bin/DNA-Pieter-64
iRep: 1.43 r^2 : 0.99 avg. cov: 22.02 % windows: 100.00



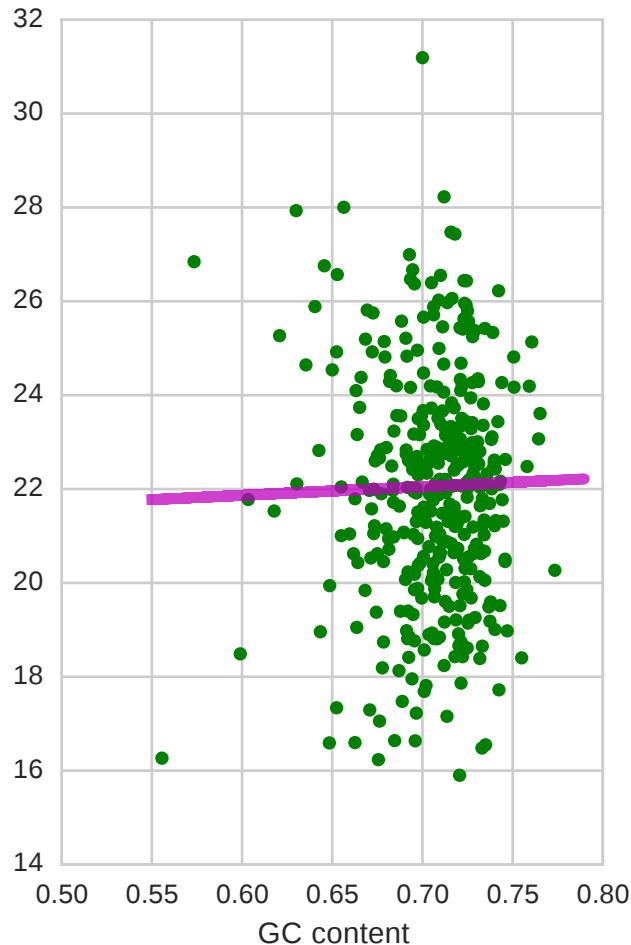
genome: ./bins/Betla_bin-contigs sample: ./splitBam/Betla_bin/DNA-Pieter-64

iRep: 1.43 raw iRep: 1.65 GC r²: 0.58 GC bias: -59.30 avg. cov: 22.02 % windows: 100.00



filtered coverage

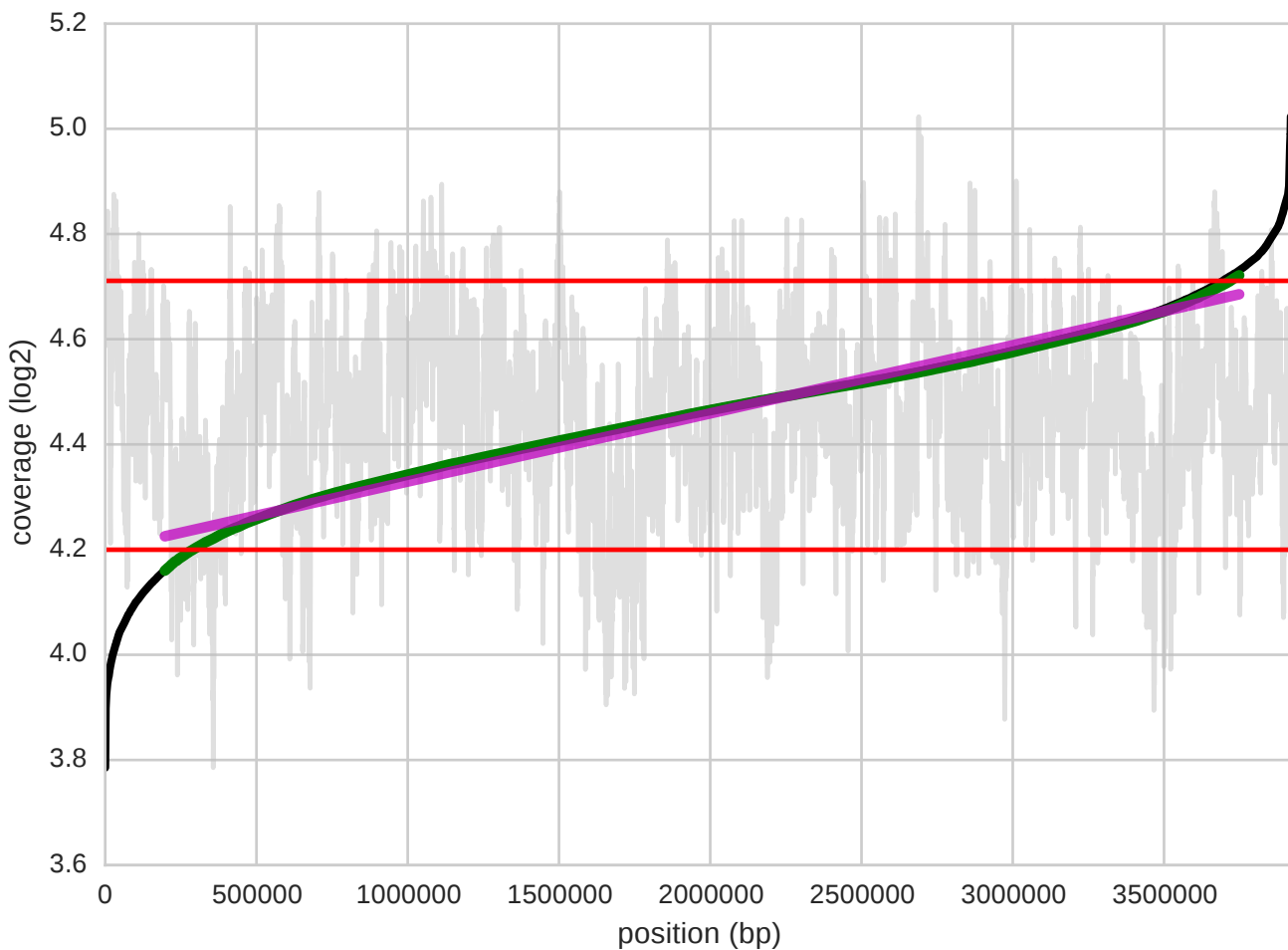
least squares fit to GC filtered data



corrected

least squares fit to corrected data

genome: ./bins/Betla_bin-contigs sample: ./splitBam/Betla_bin/DNA-Pieter-64
iRep: 1.43 un-filtered iRep: 1.43 r^2 : 0.99 avg. cov: 22.02 % windows: 100.00



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