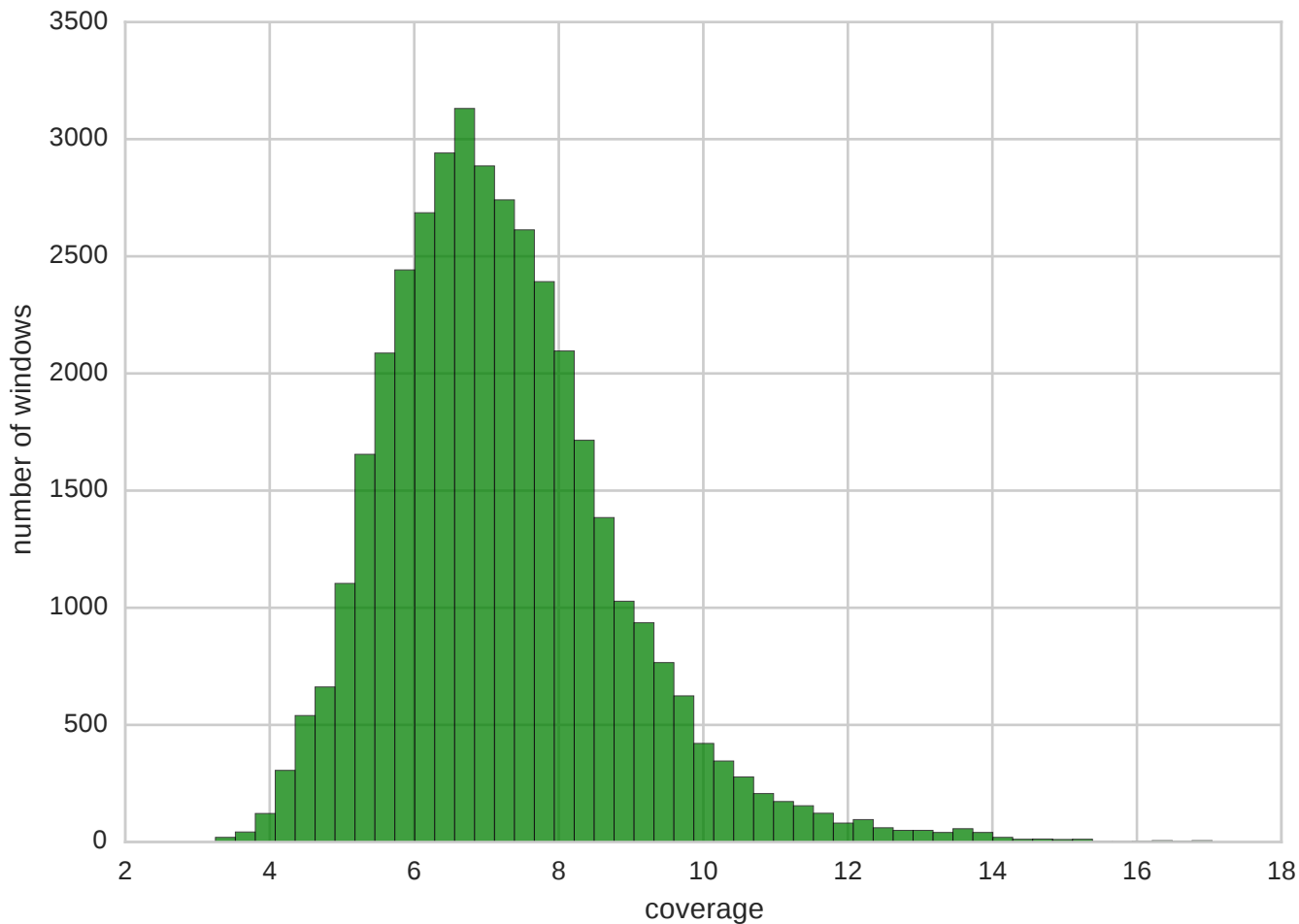
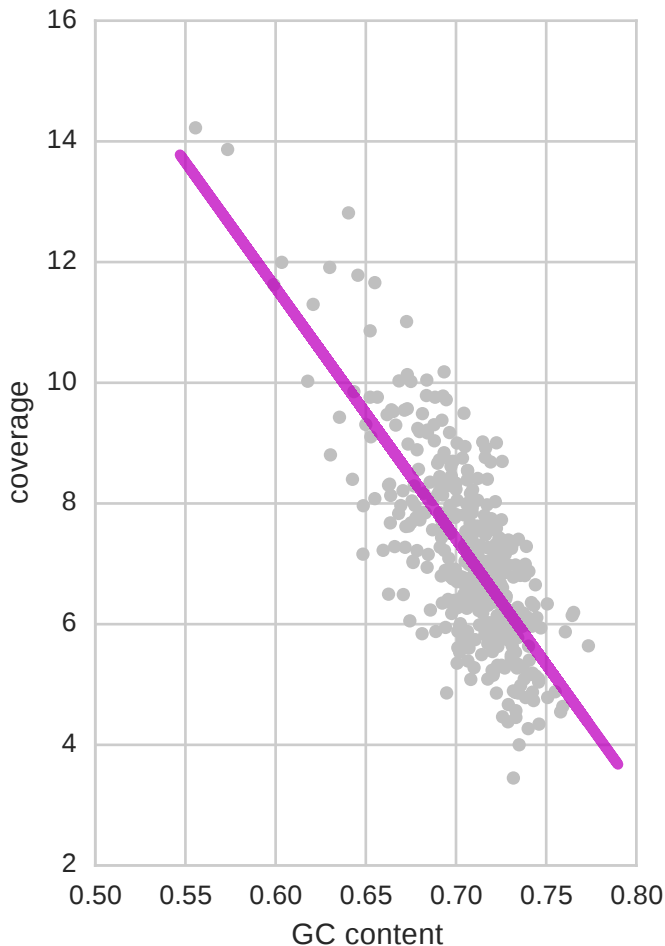


genome: ./bins/Betla_bin-contigs sample: ./splitBam/Betla_bin/DNA-Pieter-16
iRep: 1.56 r^2 : 0.99 avg. cov: 7.20 % windows: 100.00



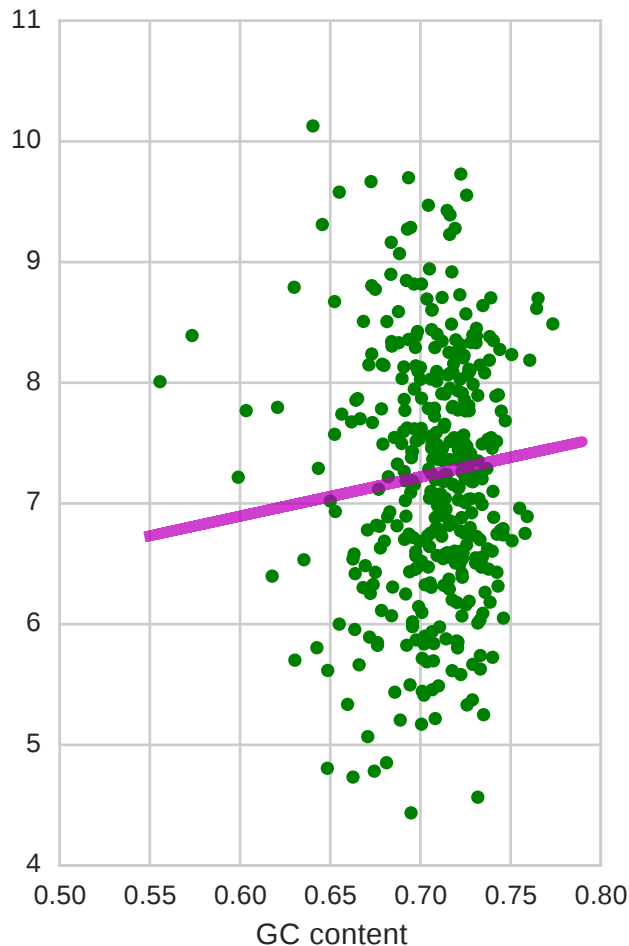
genome: ./bins/Betla_bin-contigs sample: ./splitBam/Betla_bin/DNA-Pieter-16

iRep: 1.56 raw iRep: 1.89 GC r²: 0.56 GC bias: -23.41 avg. cov: 7.20 % windows: 100.00



filtered coverage

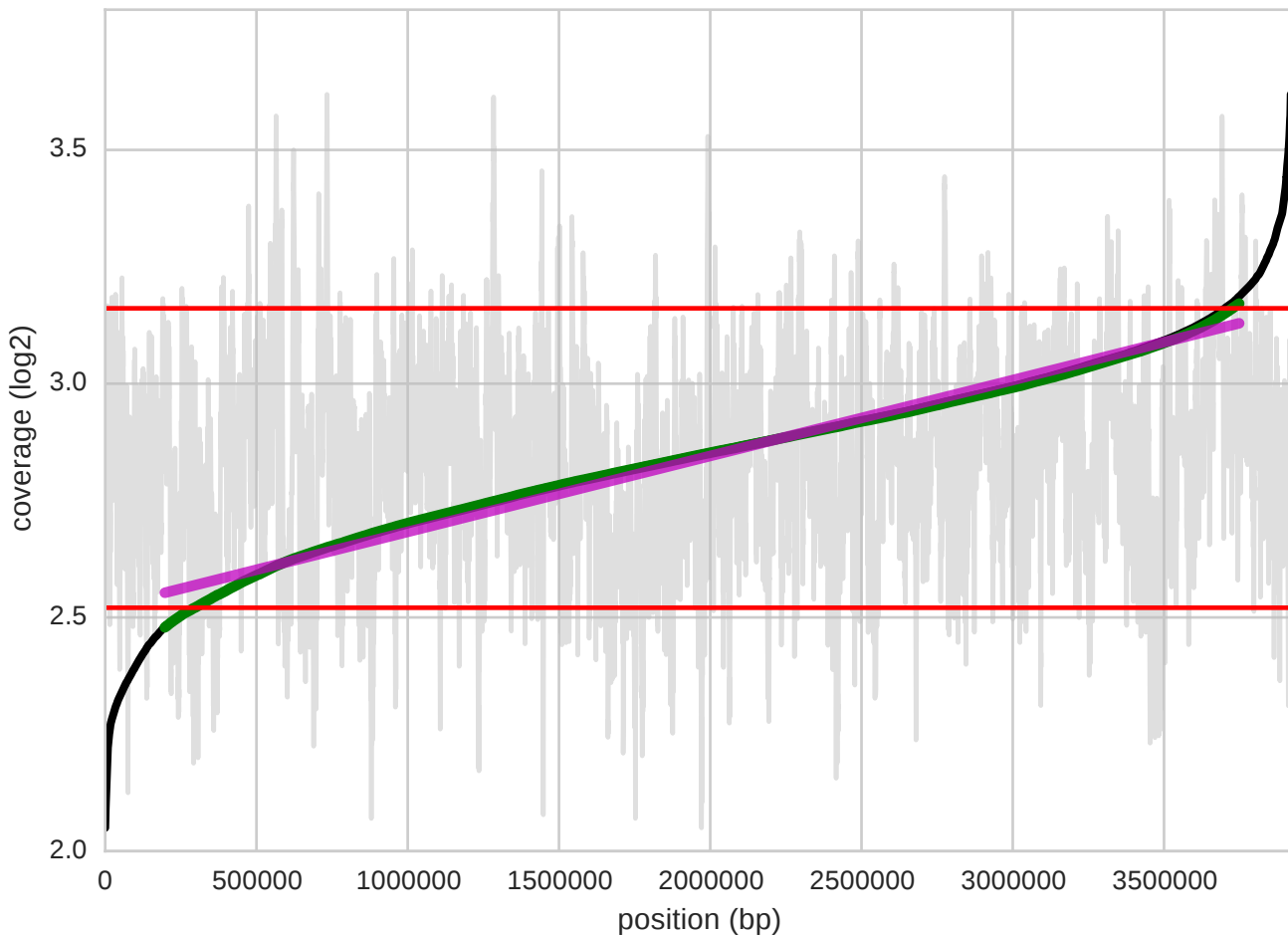
least squares fit to GC filtered data



corrected

least squares fit to corrected data

genome: ./bins/Betla_bin-contigs sample: ./splitBam/Betla_bin/DNA-Pieter-16
iRep: 1.56 un-filtered iRep: 1.56 r^2 : 0.99 avg. cov: 7.20 % windows: 100.00



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