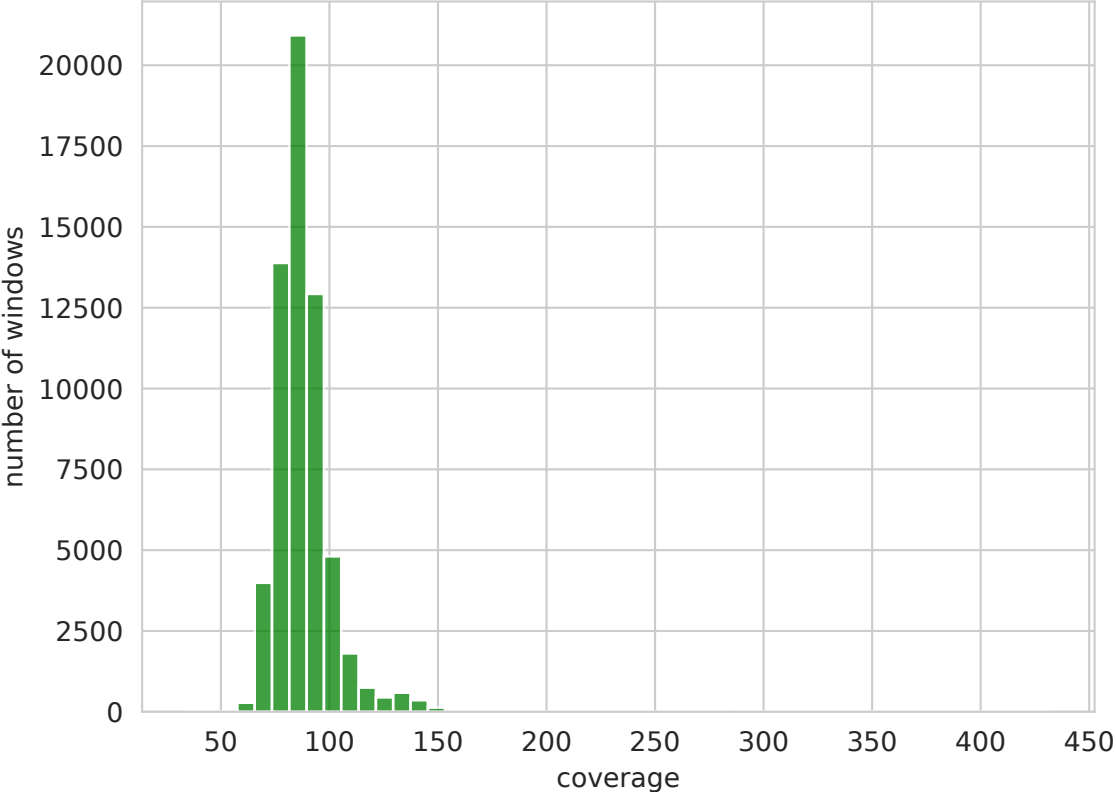
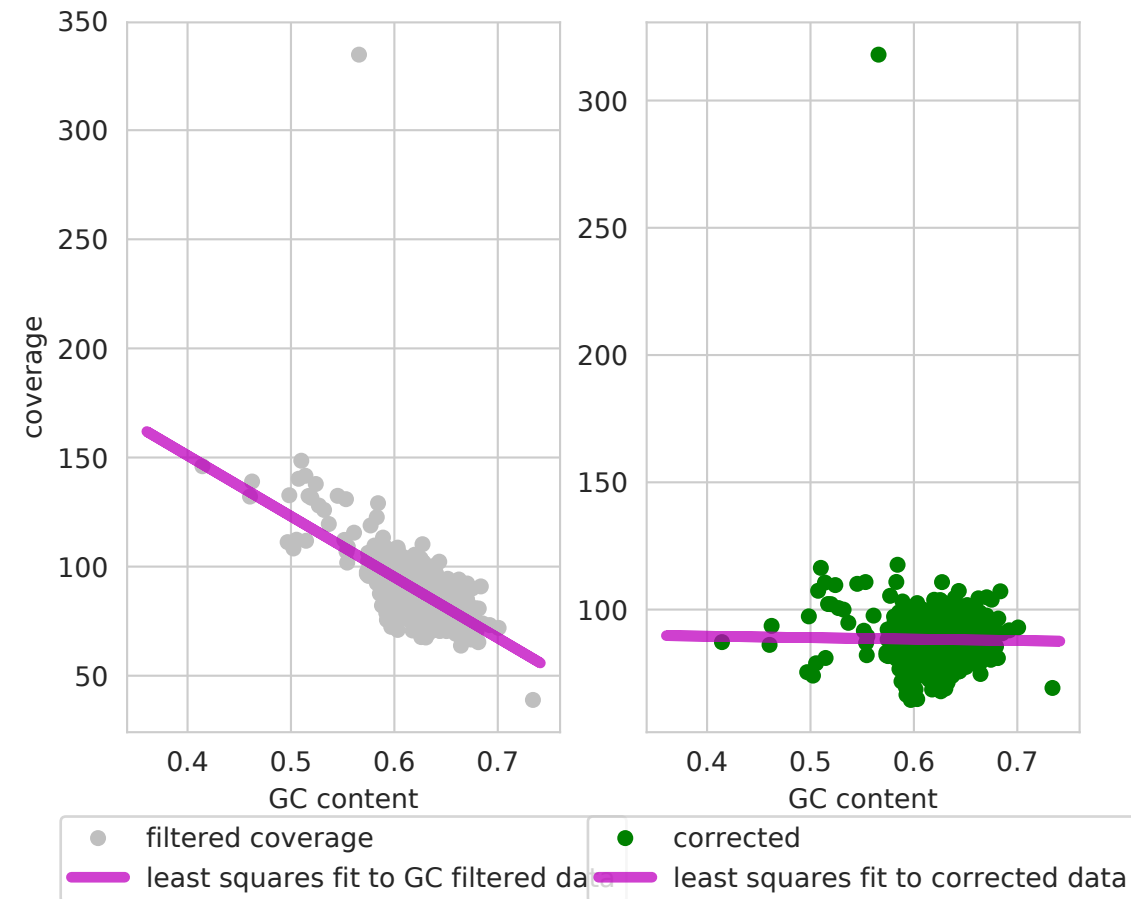


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0711/GCF_005937955.1_ASM593795v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0711-K
iRep: 1.30 r²: 0.99 avg. cov: 88.29 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0711/GCF_005937955.1_ASM593795v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0711-K
iRep: 1.30 raw iRep: 1.41 GC r^2 : 0.58 GC bias: -162.93 avg. cov: 88.29 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0711/GCF_005937955.1_ASM593795v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0711-K
iRep: 1.30 un-filtered iRep: 1.30 r^2 : 0.99 avg. cov: 88.29 % windows: 100.00

