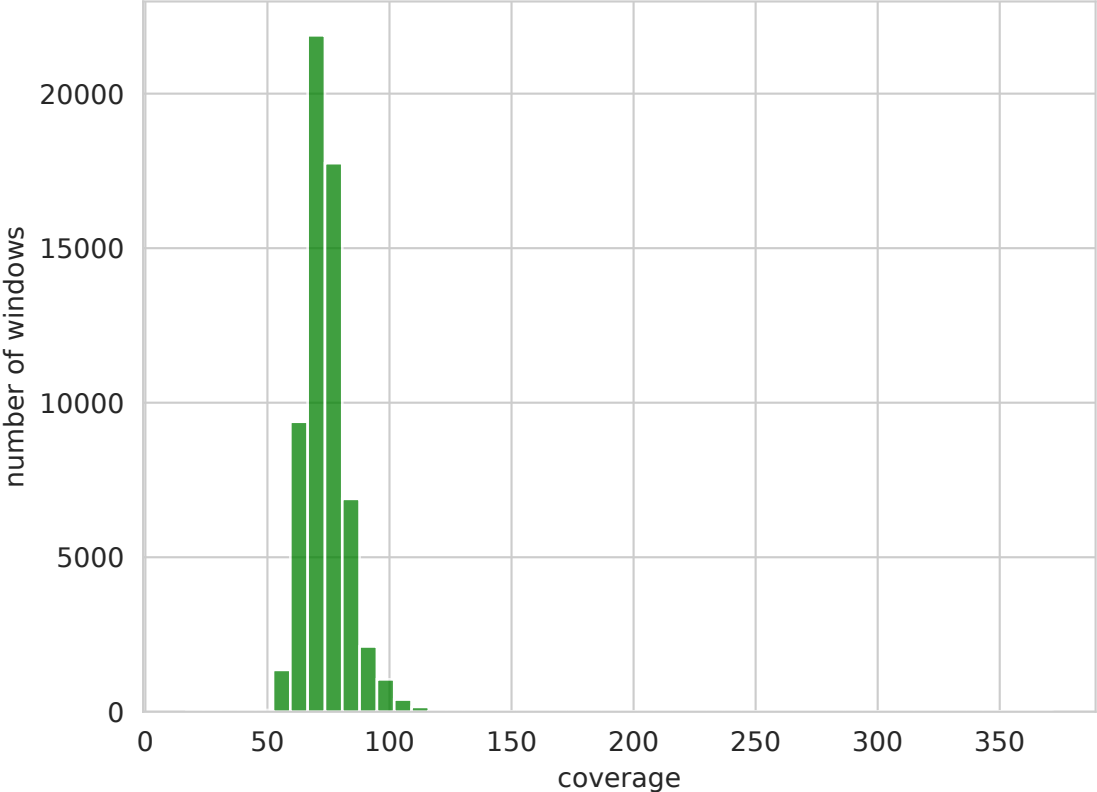
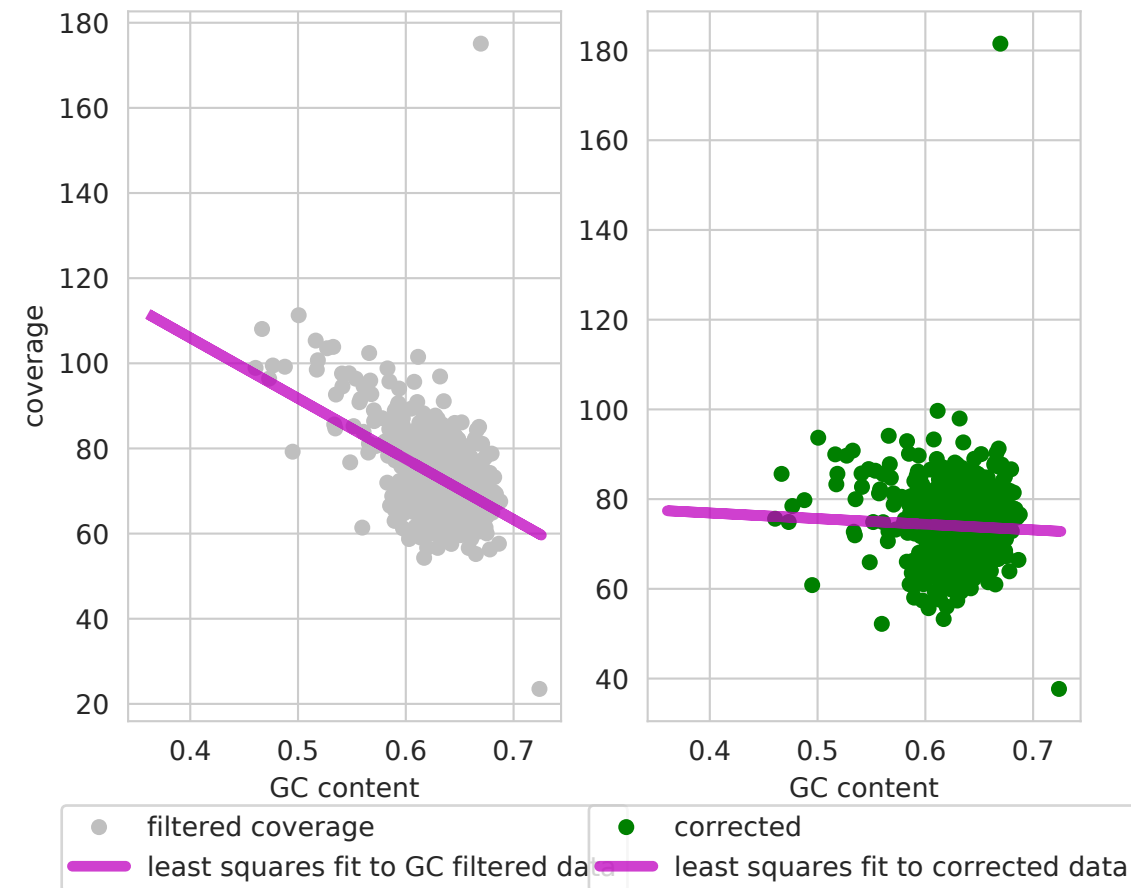


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0711/GCF_005937955_2_ASM593795v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0711-C
iRep: 1.33 r²: 0.99 avg. cov: 74.06 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0711/GCF_005937955_2_ASM593795v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0711-C
iRep: 1.33 raw iRep: 1.38 GC r^2 : 0.32 GC bias: -45.55 avg. cov: 74.06 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0711/GCF_005937955.2_ASM593795v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0711-C
iRep: 1.33 un-filtered iRep: 1.33 r^2 : 0.99 avg. cov: 74.06 % windows: 100.00

