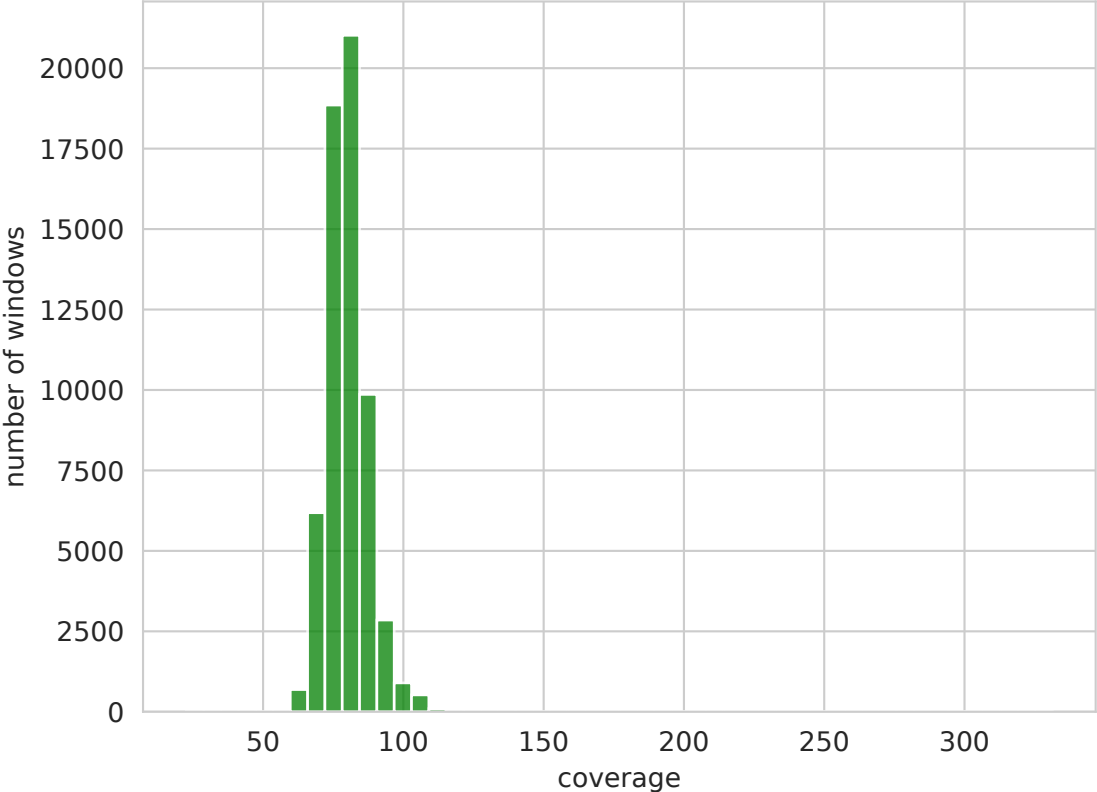
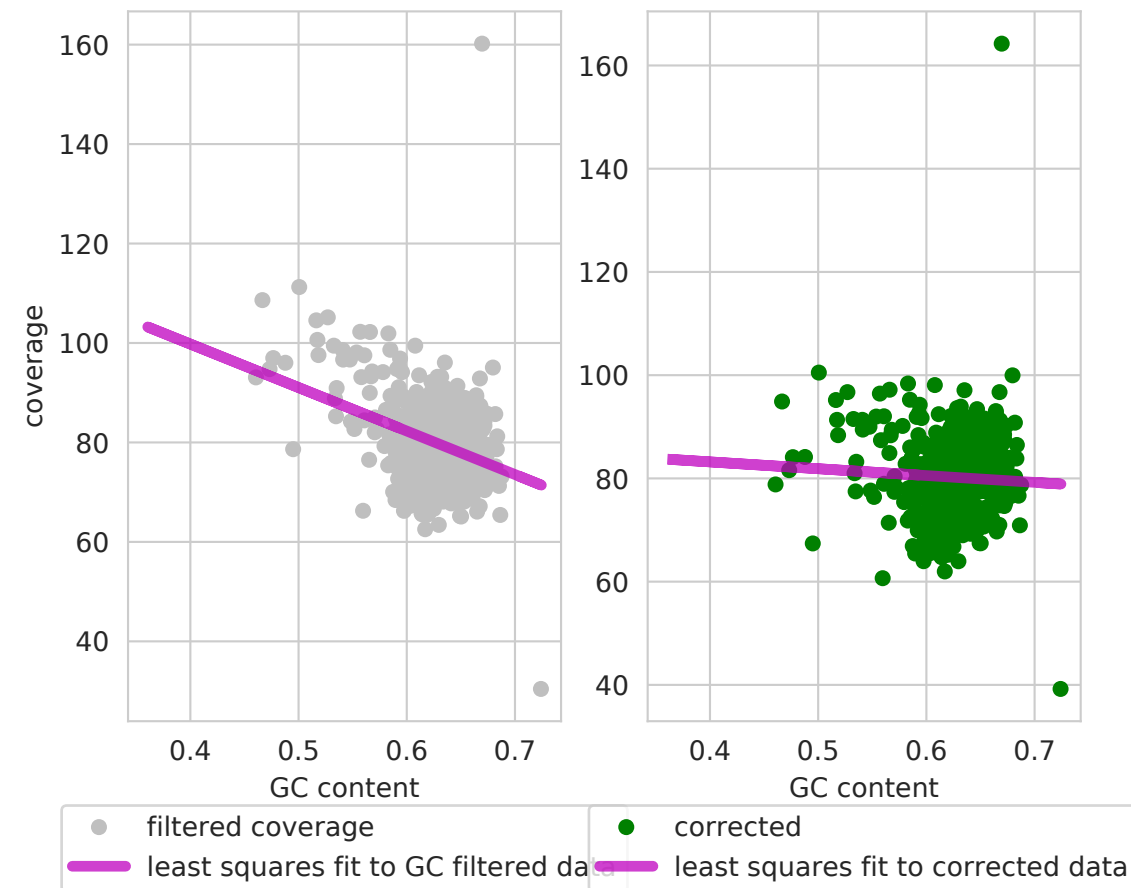


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0711/GCF_005937955_2_ASM593795v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0711-A
iRep: 1.28 r^2 : 0.99 avg. cov: 80.22 % windows: 100.00



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-A
iRep: 1.28 raw iRep: 1.29 GC r^2 : 0.17 GC bias: -14.99 avg. cov: 80.22 % 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-A
iRep: 1.28 un-filtered iRep: 1.28 r^2 : 0.99 avg. cov: 80.22 % windows: 100.00

