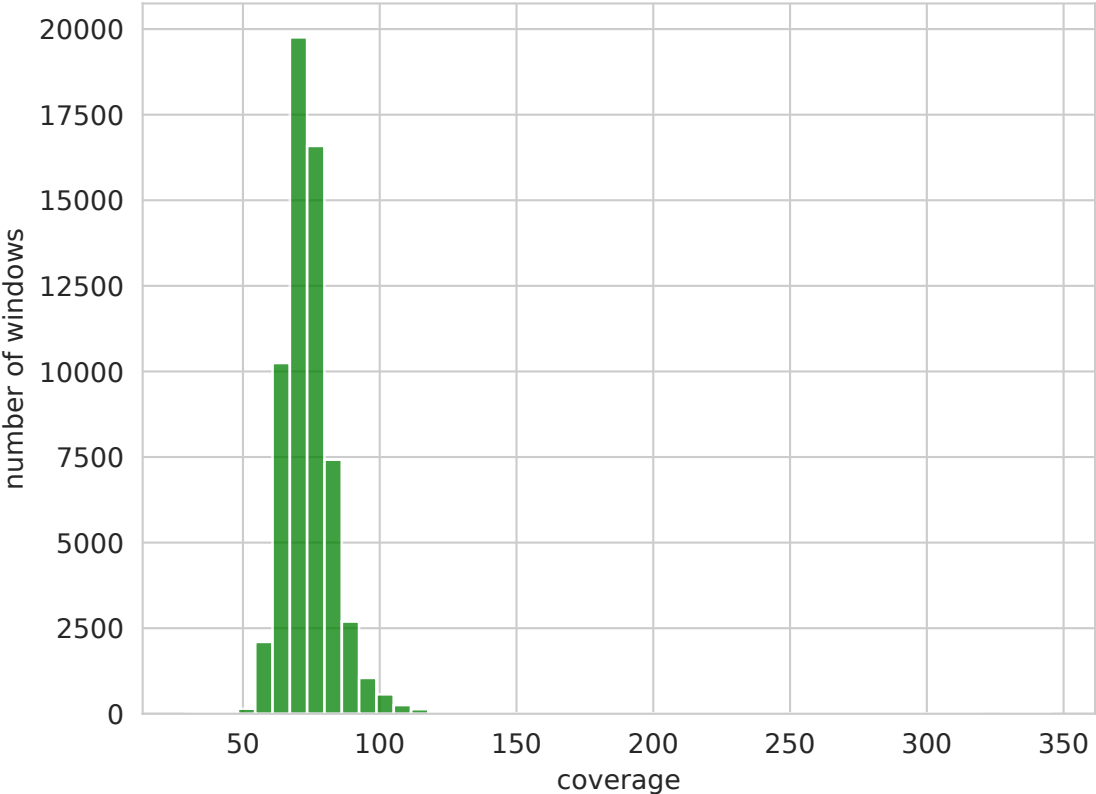
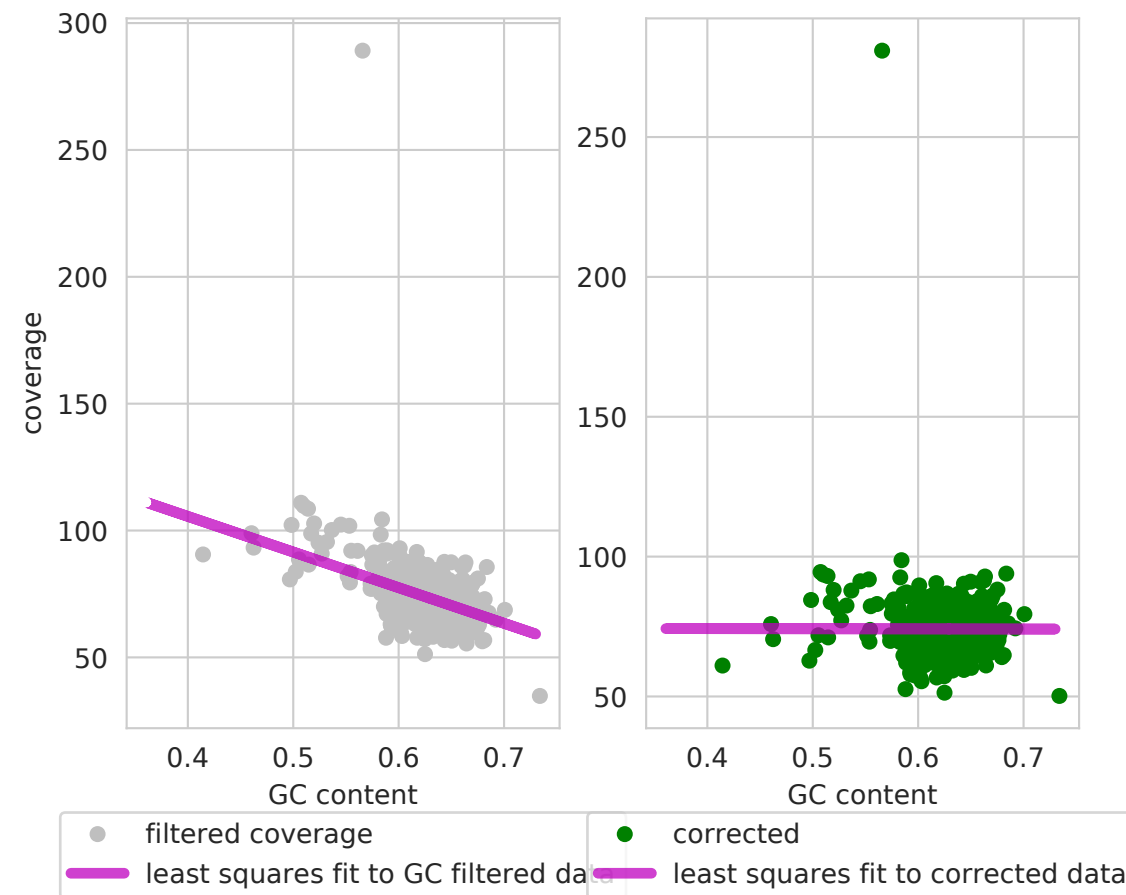


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0711/GCF\_005937955.1\_ASM593795v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0711-C  
iRep: 1.34     $r^2$ : 0.99    avg. cov: 74.16    % windows: 100.00



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-C  
iRep: 1.34 raw iRep: 1.38 GC  $r^2$ : 0.31 GC bias: -43.31 avg. cov: 74.16 % 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-C  
iRep: 1.34 un-filtered iRep: 1.34  $r^2$ : 0.99 avg. cov: 74.16 % windows: 100.00

