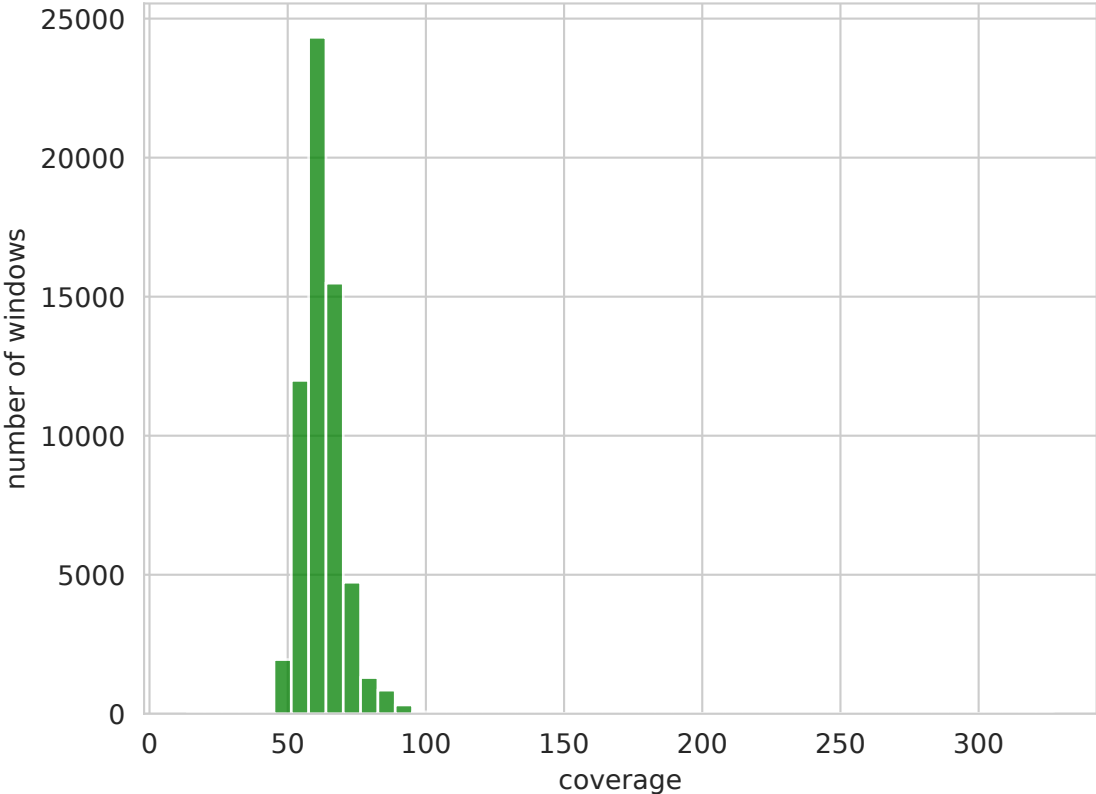
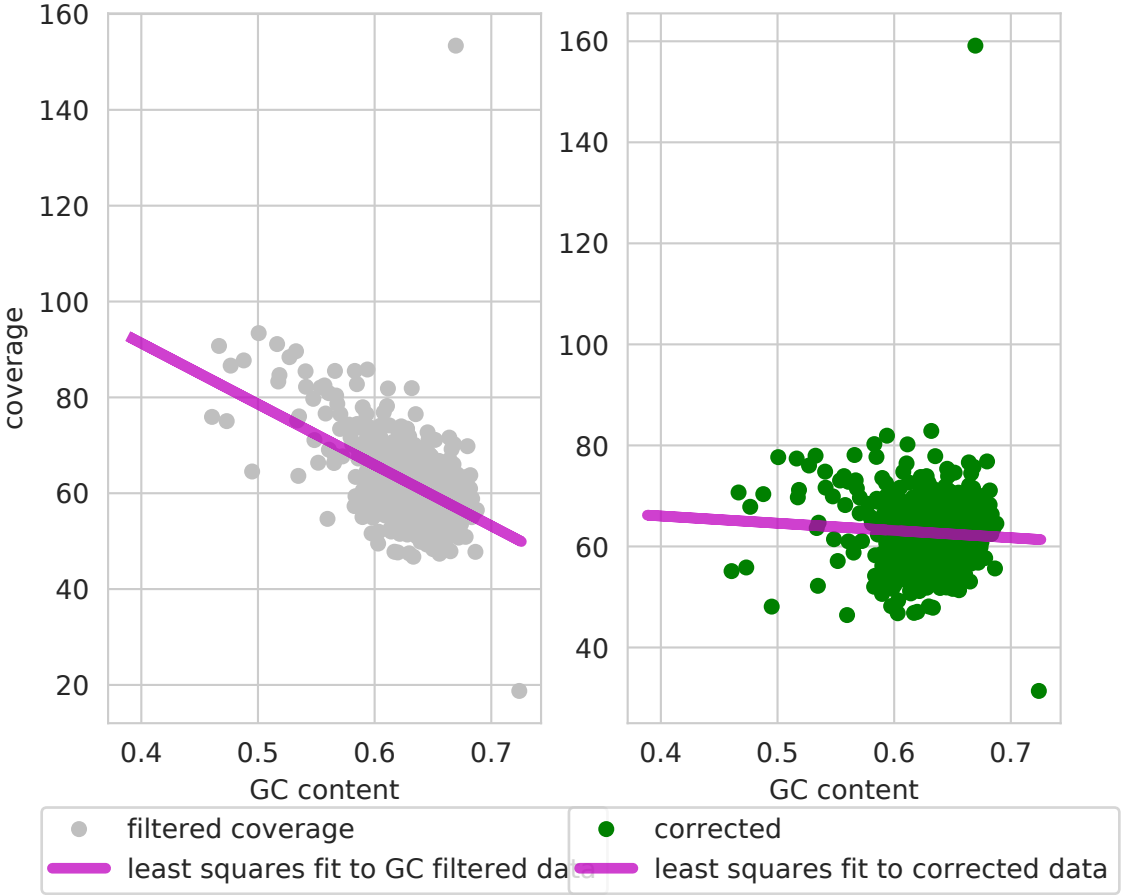


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0711/GCF\_005937955\_2\_ASM593795v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0711-D  
iRep: 1.32    r<sup>2</sup>: 0.99    avg. cov: 62.78    % windows: 100.00



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-D  
iRep: 1.32 raw iRep: 1.36 GC  $r^2$ : 0.35 GC bias: -44.33 avg. cov: 62.78 % 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-D  
iRep: 1.32 un-filtered iRep: 1.32  $r^2$ : 0.99 avg. cov: 62.78 % windows: 100.00

