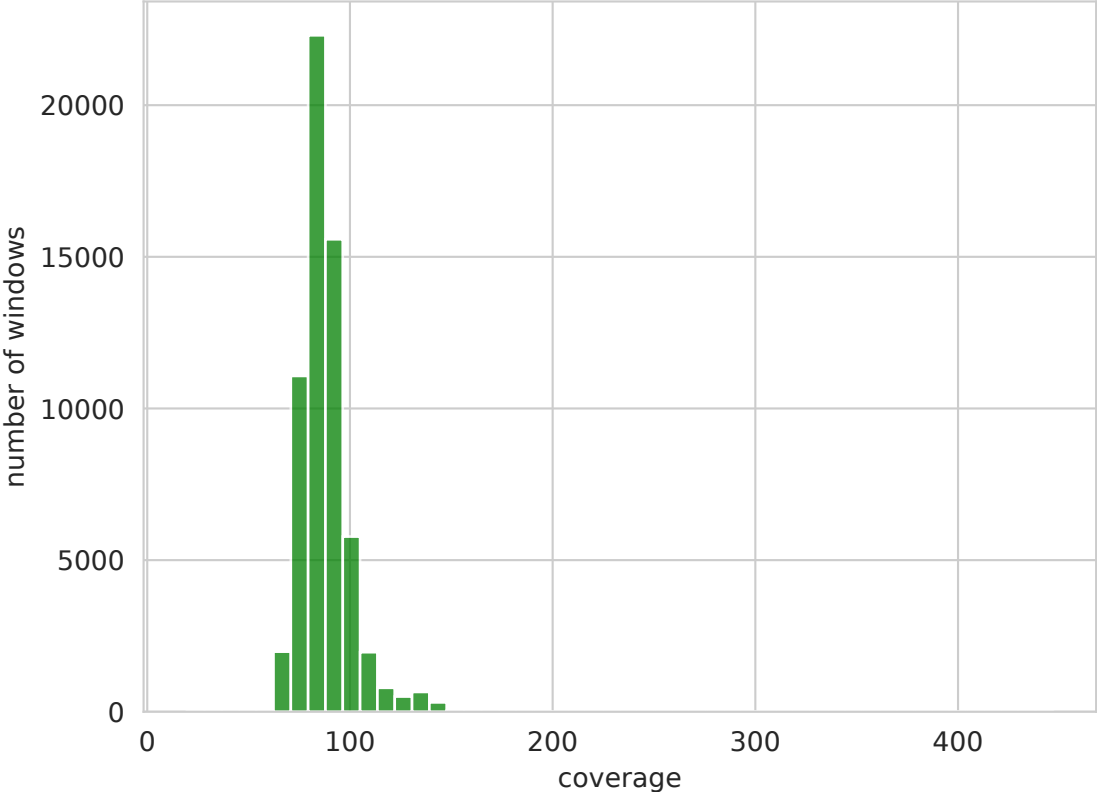
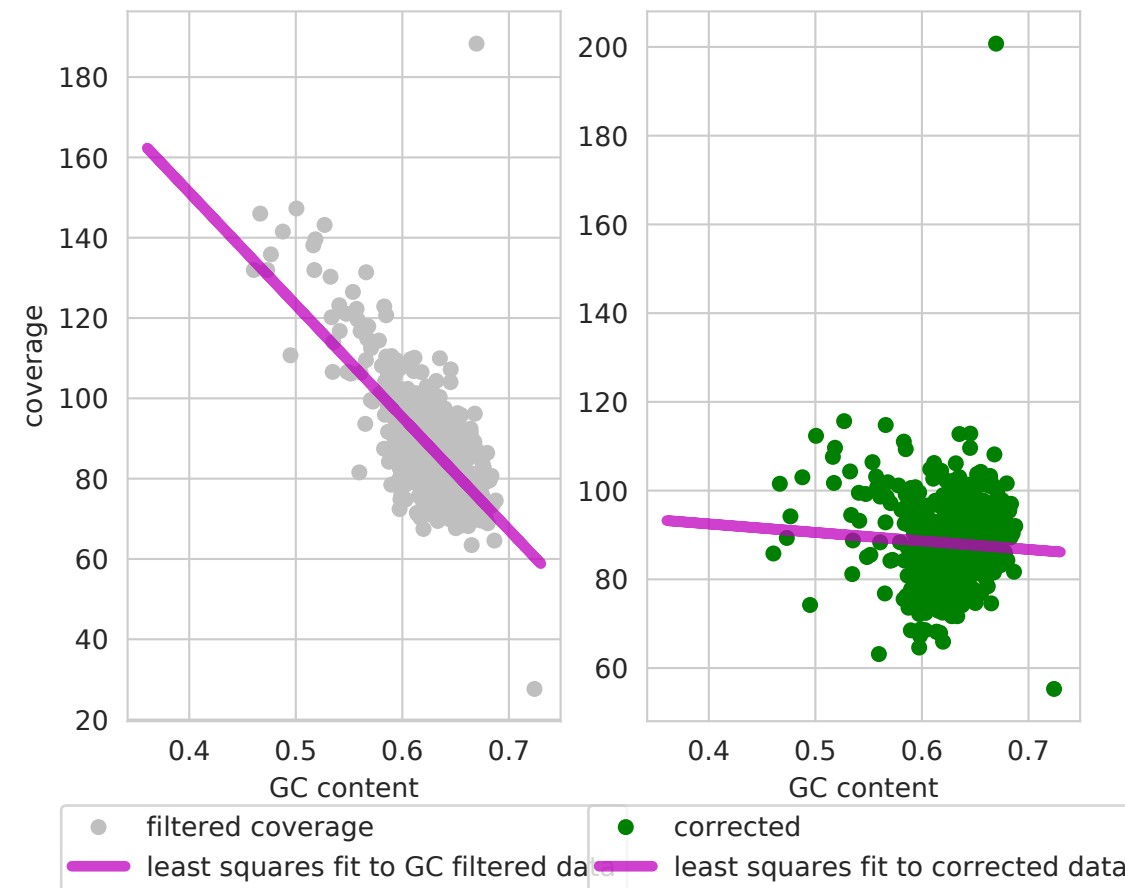


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0711/GCF\_005937955\_2\_ASM593795v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0711-K  
iRep: 1.30    r<sup>2</sup>: 0.99    avg. cov: 88.16    % windows: 100.00



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-K  
iRep: 1.30 raw iRep: 1.41 GC  $r^2$ : 0.59 GC bias: -165.84 avg. cov: 88.16 % 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-K  
iRep: 1.30 un-filtered iRep: 1.30  $r^2$ : 0.99 avg. cov: 88.16 % windows: 100.00

