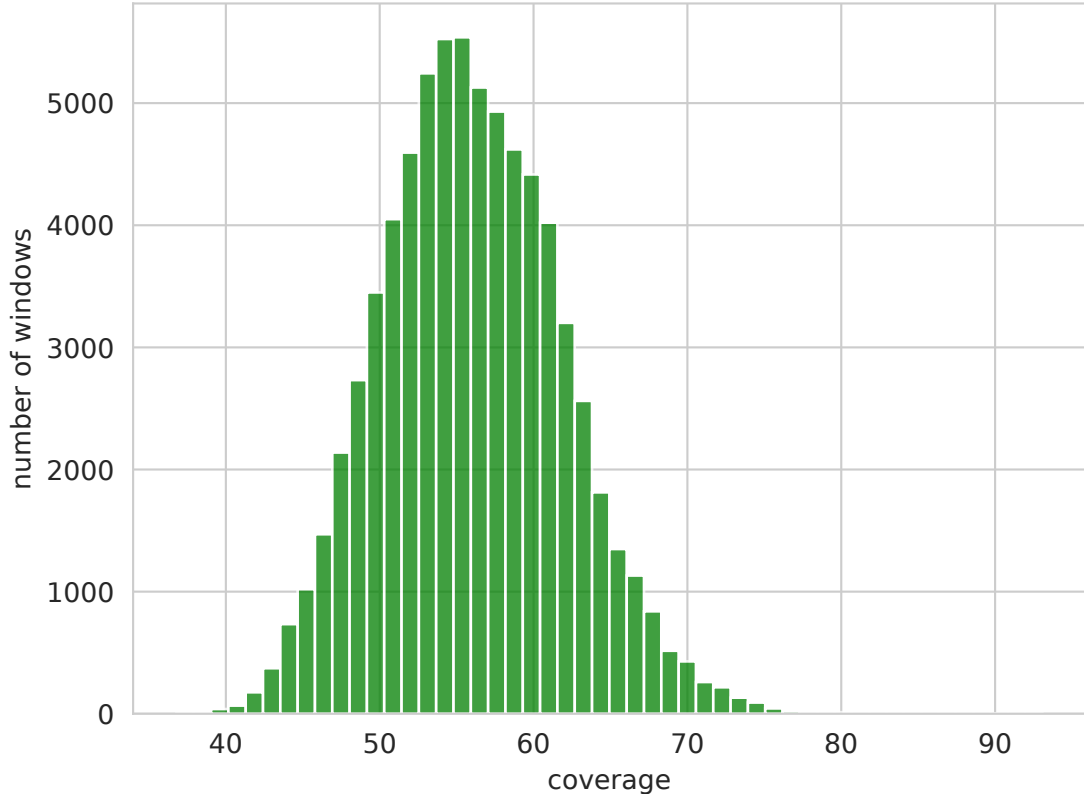
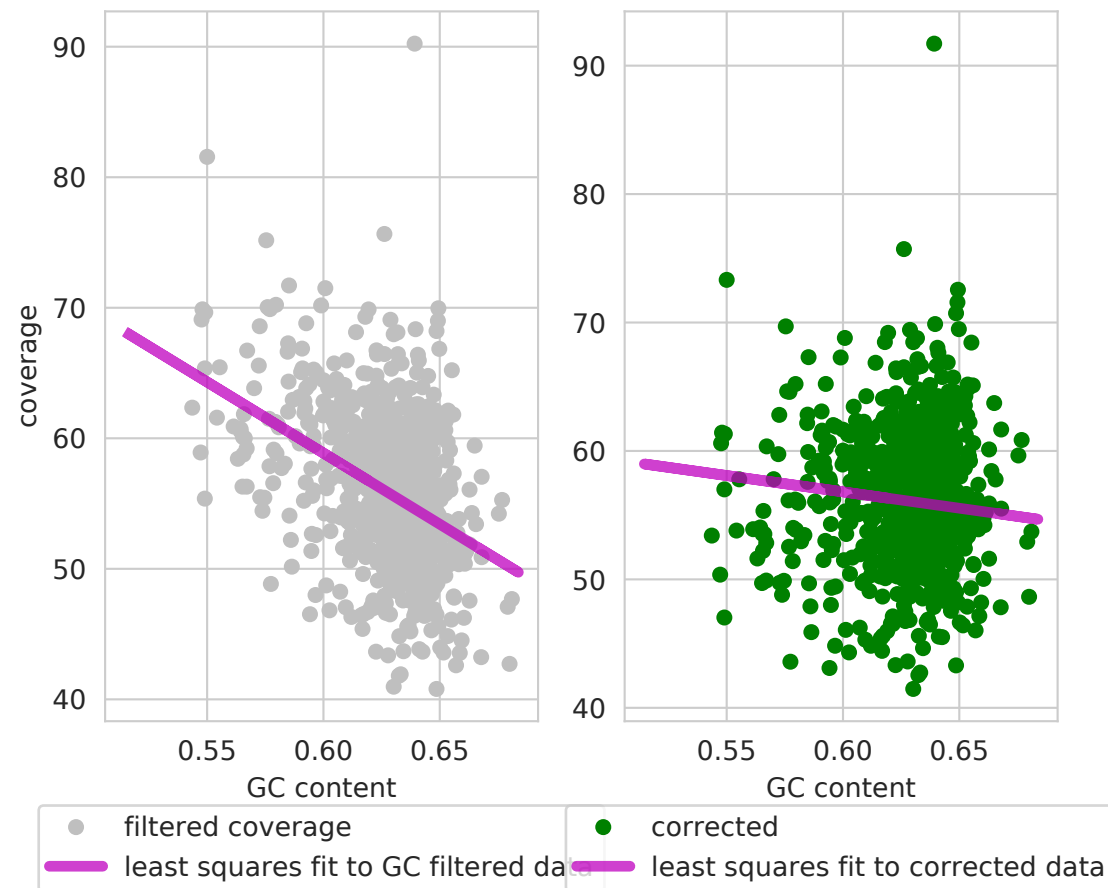


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0727/GCF\_005937885\_2\_ASM593788v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0727-C  
iRep: 1.34     $r^2$ : 0.99    avg. cov: 56.08    % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0727/GCF\_005937885\_2\_ASM593788v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0727-C  
iRep: 1.34 raw iRep: 1.38 GC  $r^2$ : 0.19 GC bias: -20.61 avg. cov: 56.08 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0727/GCF\_005937885\_2\_ASM593788v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0727-C  
iRep: 1.34   un-filtered iRep: 1.34    $r^2$ : 0.99   avg. cov: 56.08   % windows: 100.00

