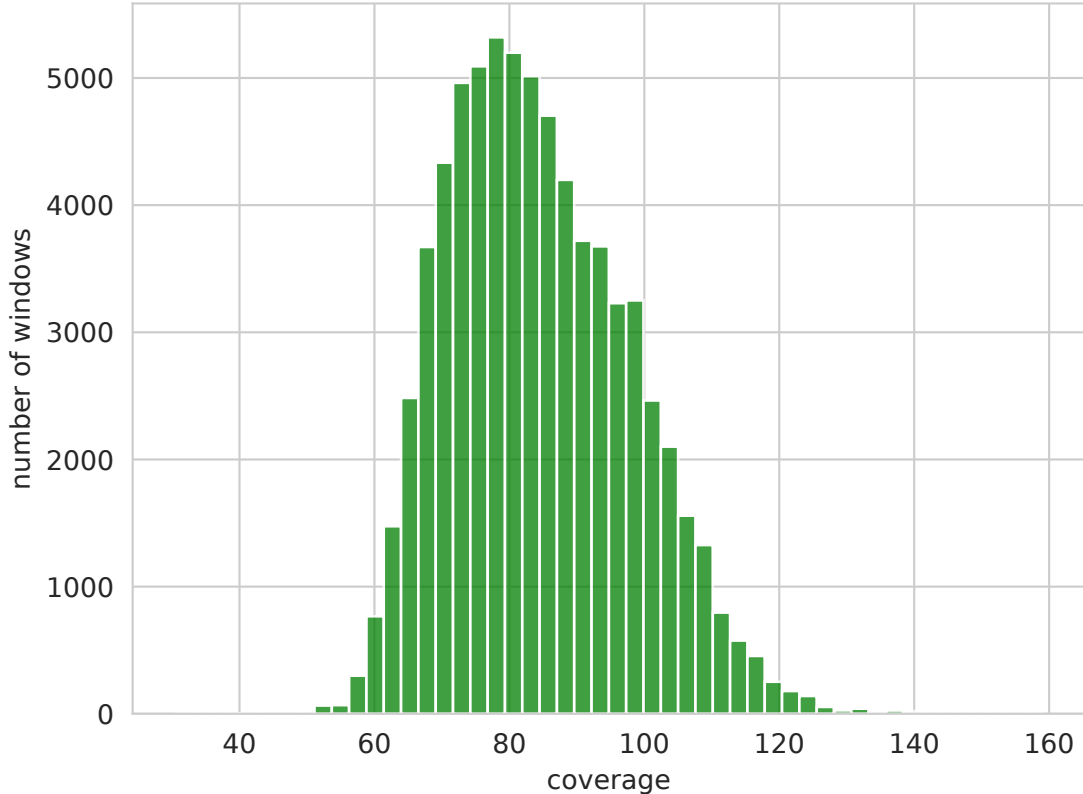
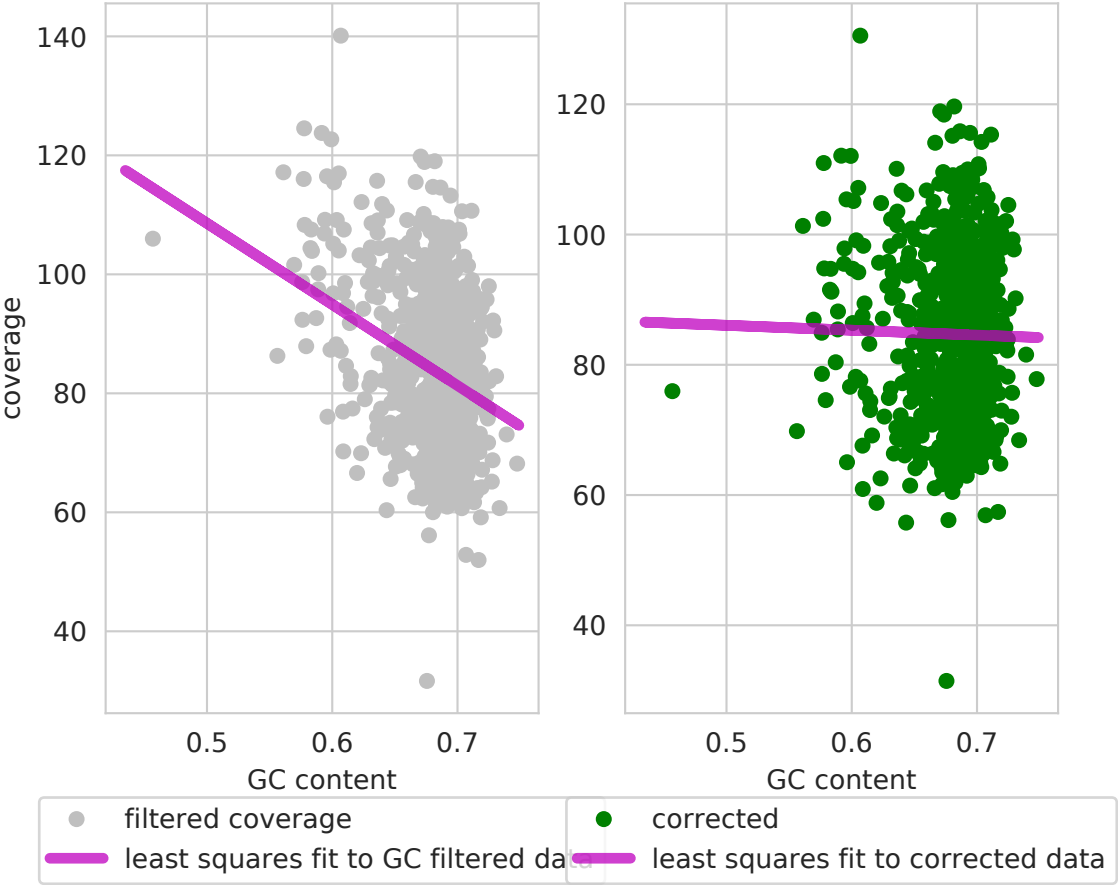


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0712/GCF\_006323185.1\_ASM632318v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0712-C  
iRep: 1.62     $r^2$ : 1.00    avg. cov: 84.43    % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0712/GCF\_006323185.1\_ASM632318v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0712-C  
iRep: 1.62 raw iRep: 1.65 GC r<sup>2</sup>: 0.10 GC bias: -14.32 avg. cov: 84.43 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0712/GCF\_006323185.1\_ASM632318v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0712-C  
iRep: 1.62 un-filtered iRep: 1.62  $r^2$ : 1.00 avg. cov: 84.43 % windows: 100.00

