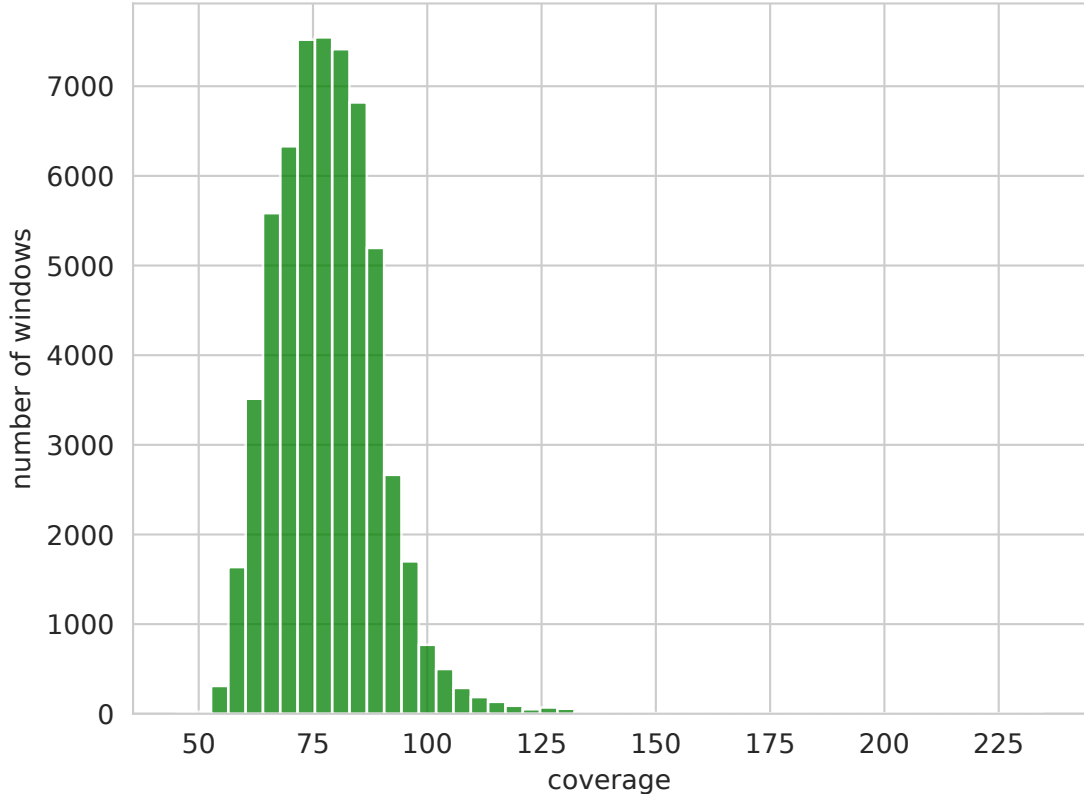
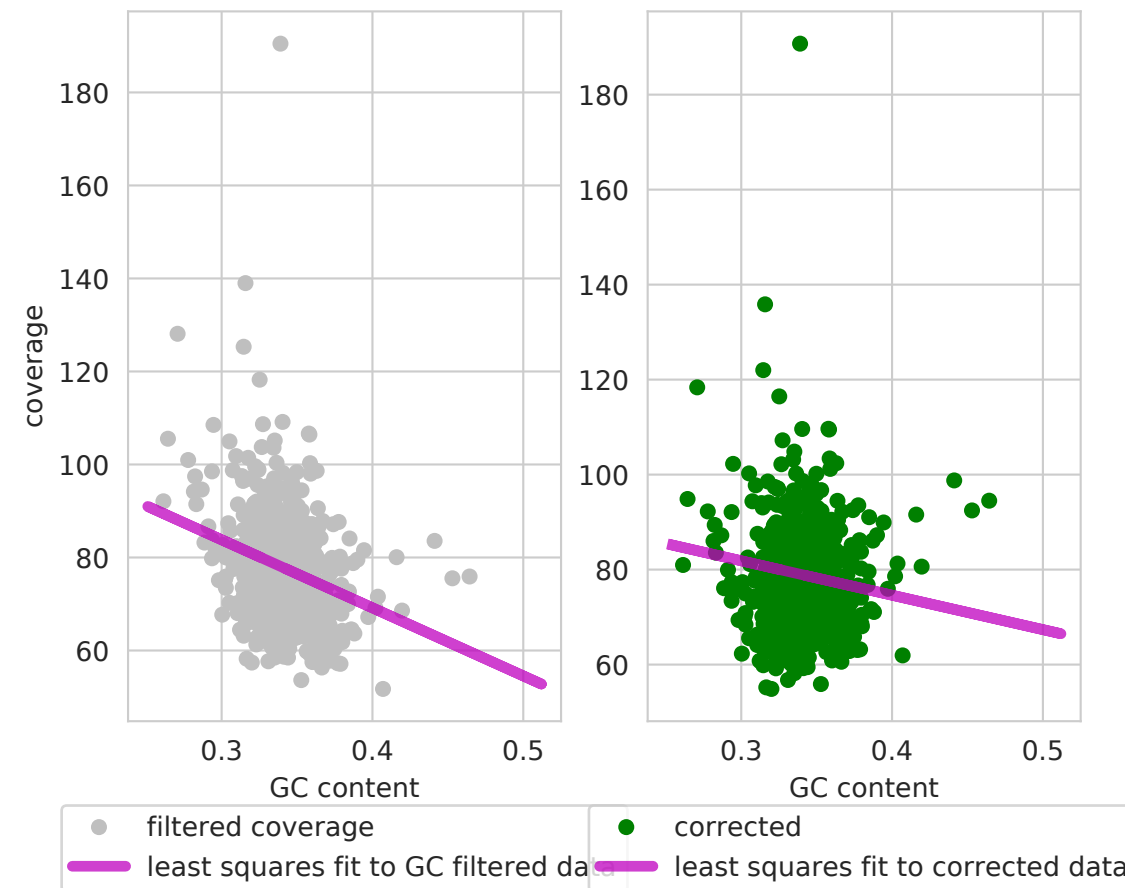


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0721/GCF\_002007065\_3\_ASM200706v3\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0721-B  
iRep: 1.49     $r^2$ : 0.99    avg. cov: 78.36    % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0721/GCF\_002007065\_3\_ASM200706v3\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0721-B  
iRep: 1.49 raw iRep: 1.53 GC  $r^2$ : 0.10 GC bias: -14.40 avg. cov: 78.36 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0721/GCF\_002007065\_3\_ASM200706v3\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0721-B  
iRep: 1.49 un-filtered iRep: 1.49  $r^2$ : 0.99 avg. cov: 78.36 % windows: 100.00

