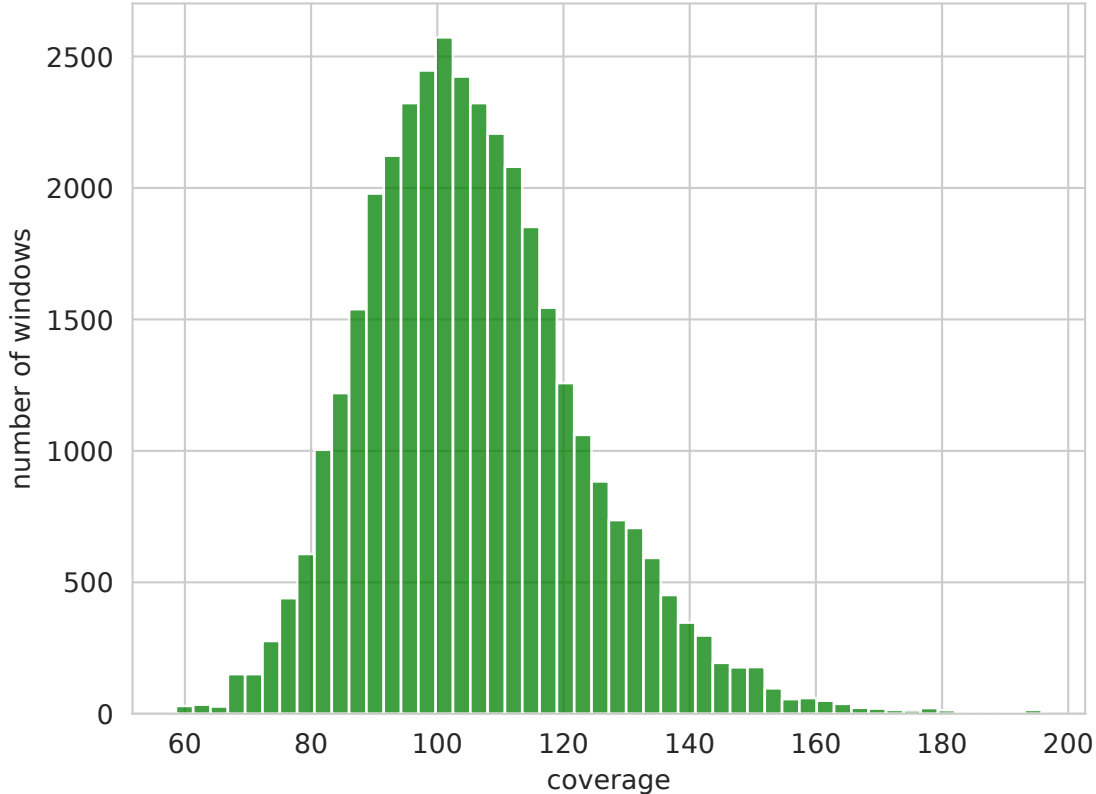
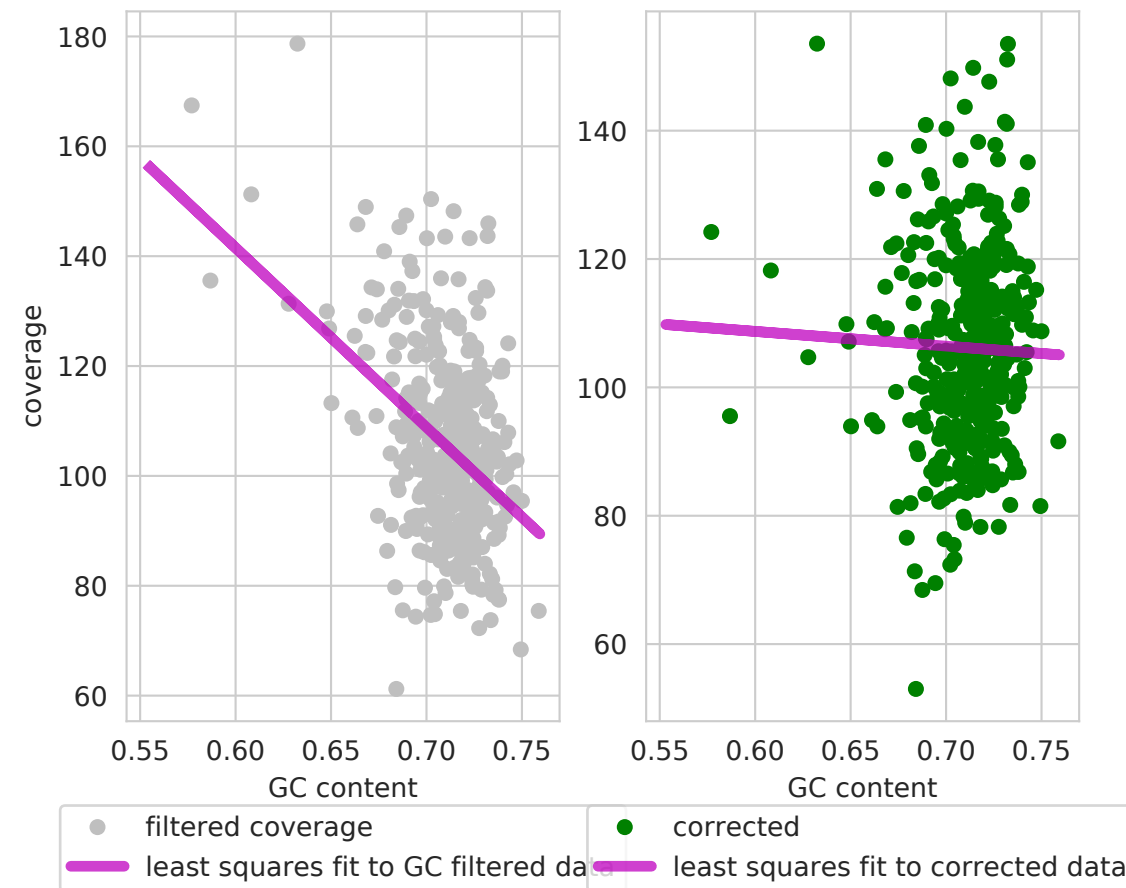


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0715/GCF\_005938005.1\_ASM593800v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0715-A  
iRep: 1.56   r<sup>2</sup>: 0.99   avg. cov: 105.76   % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0715/GCF\_005938005.1\_ASM593800v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0715-A  
iRep: 1.56 raw iRep: 1.61 GC r<sup>2</sup>: 0.19 GC bias: -60.72 avg. cov: 105.76 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0715/GCF\_005938005.1\_ASM593800v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0715-A  
iRep: 1.56 un-filtered iRep: 1.56  $r^2$ : 0.99 avg. cov: 105.76 % windows: 100.00

