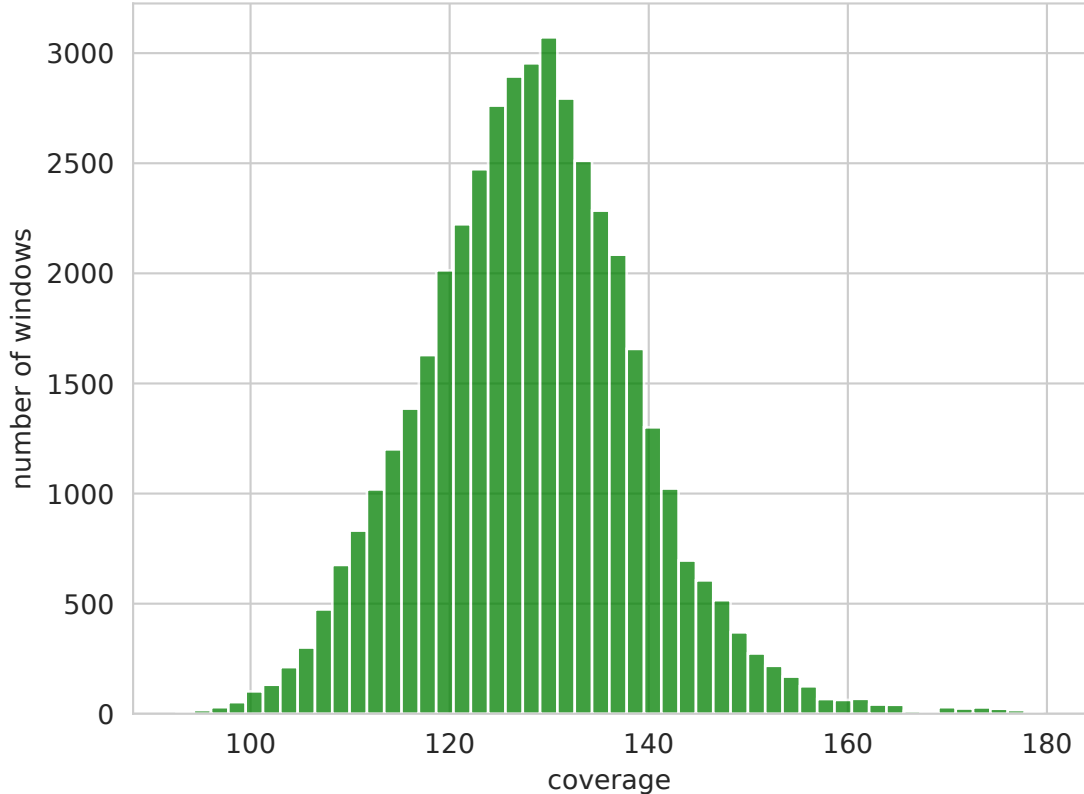
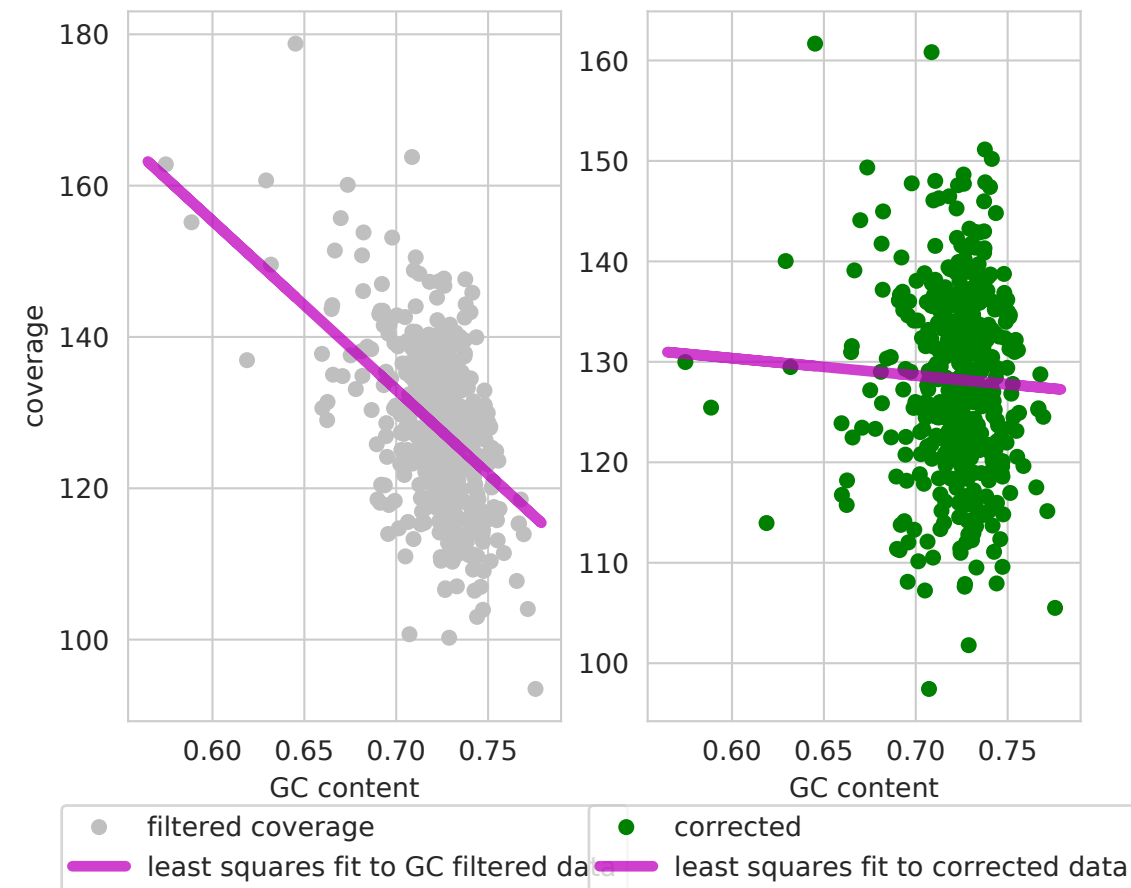


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0722/GCF\_005937995\_2\_ASM593799v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0722-C  
iRep: 1.25   r<sup>2</sup>: 0.99   avg. cov: 128.19   % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0722/GCF\_005937995\_2\_ASM593799v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0722-C  
iRep: 1.25 raw iRep: 1.28 GC  $r^2$ : 0.23 GC bias: -50.42 avg. cov: 128.19 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0722/GCF\_005937995.2\_ASM593799v2\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0722-C  
iRep: 1.25 un-filtered iRep: 1.25 r^2: 0.99 avg. cov: 128.19 % windows: 100.00

