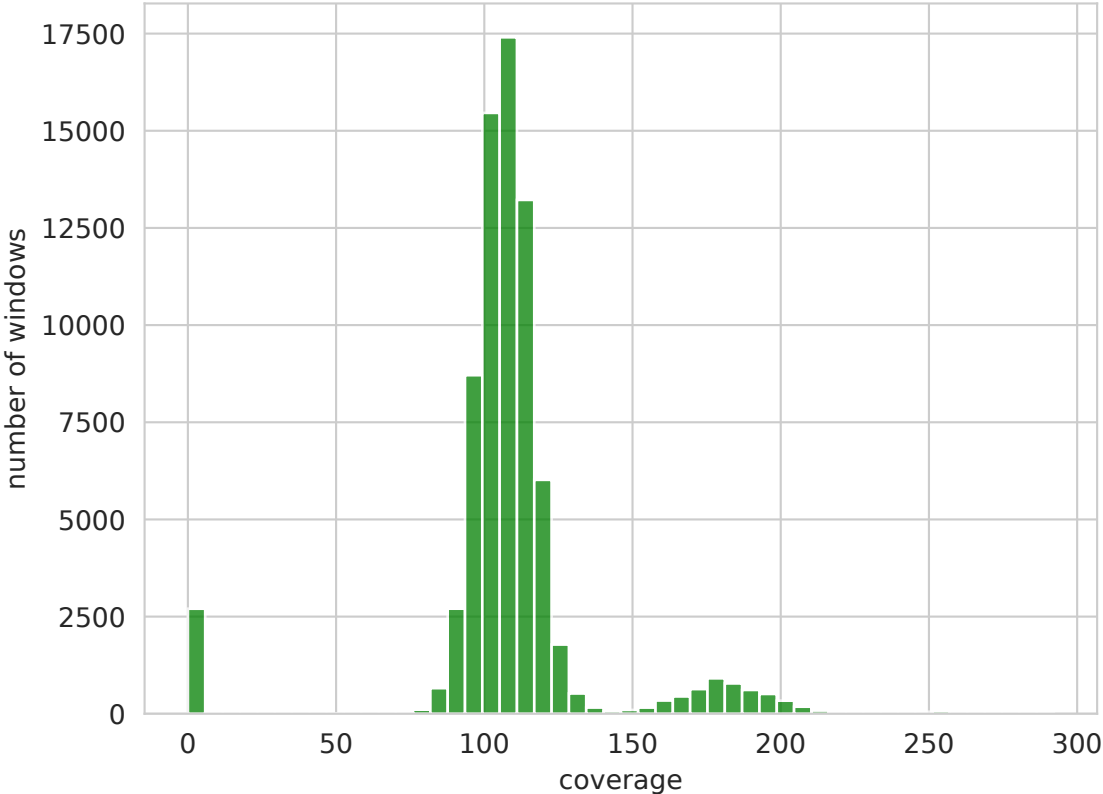
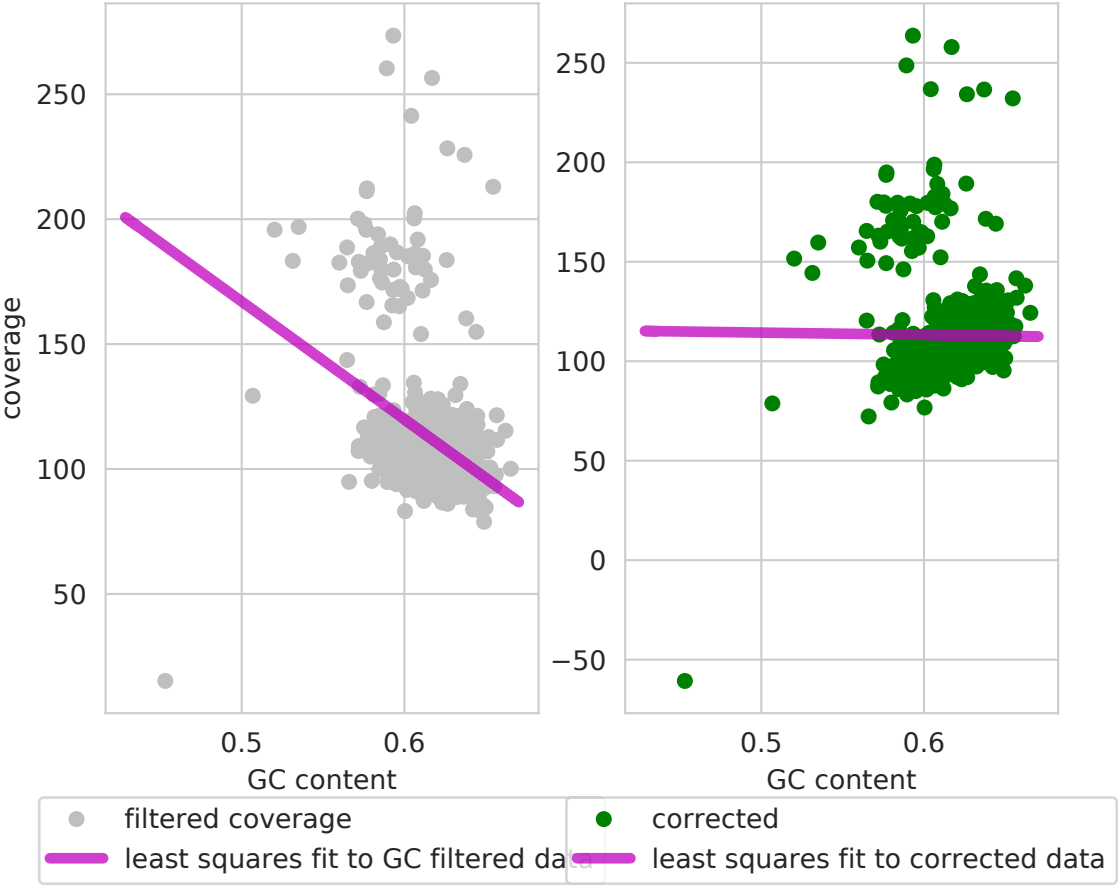


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0724/GCF\_005938745.1\_ASM593874v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0724-B  
iRep: n/a     $r^2$ : 0.86    avg. cov: 109.19    % windows: 96.38



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0724/GCF\_005938745.1\_ASM593874v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0724-B  
iRep: n/a    raw iRep: 1.41    GC r<sup>2</sup>: 0.20    GC bias: -92.09    avg. cov: 109.19    % windows: 96.38



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0724/GCF\_005938745.1\_ASM593874v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0724-B  
iRep: n/a un-filtered iRep: 1.42 r^2: 0.86 avg. cov: 109.19 % windows: 96.38

