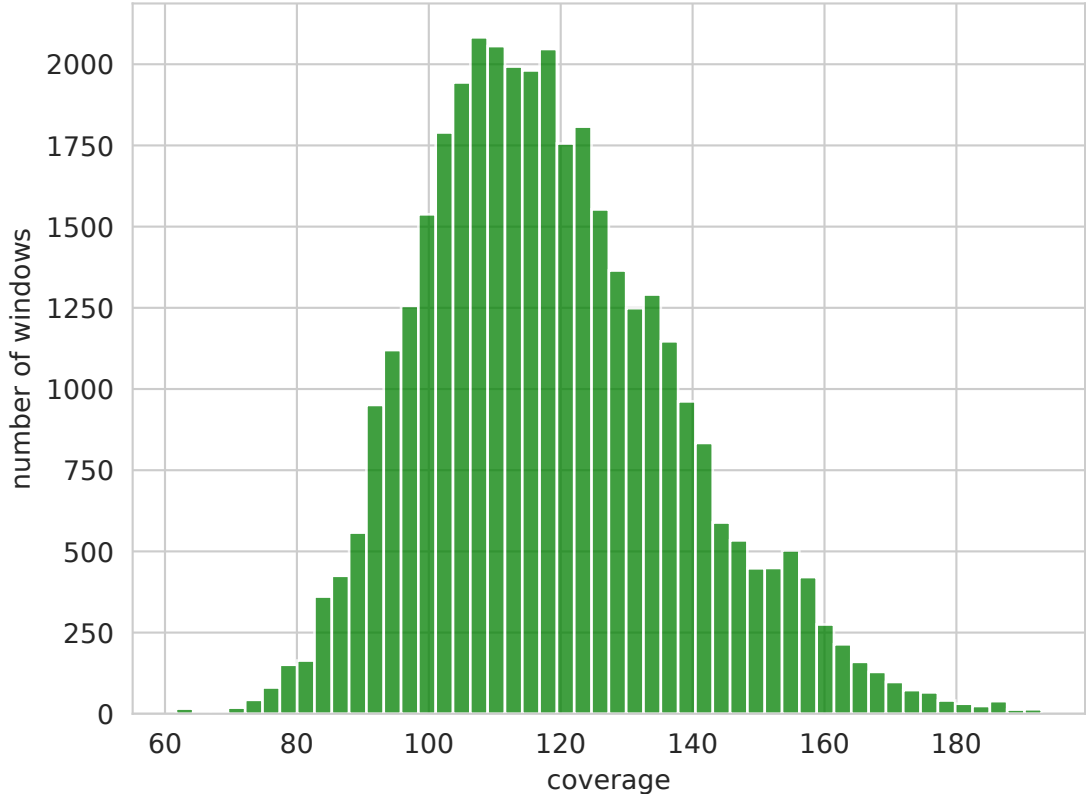
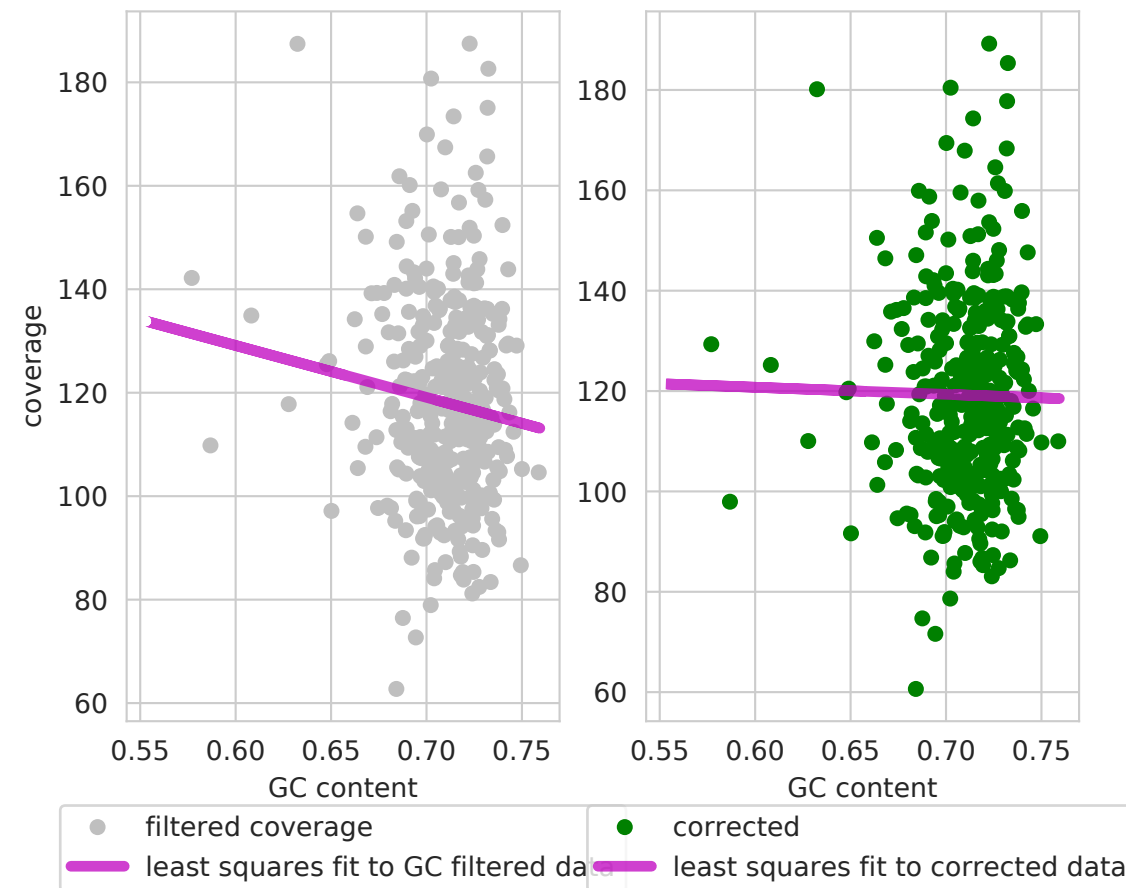


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes\_ncbi/KBS0715/GCF\_005938005.1\_ASM593800v1\_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa\_bam/KBS0715-C  
iRep: 1.64    r^2: 0.99    avg. cov: 118.54    % windows: 100.00



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-C  
iRep: 1.64 raw iRep: 1.64 GC  $r^2$ : 0.01 GC bias: -1.40 avg. cov: 118.54 % 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-C  
iRep: 1.64 un-filtered iRep: 1.64  $r^2$ : 0.99 avg. cov: 118.54 % windows: 100.00

