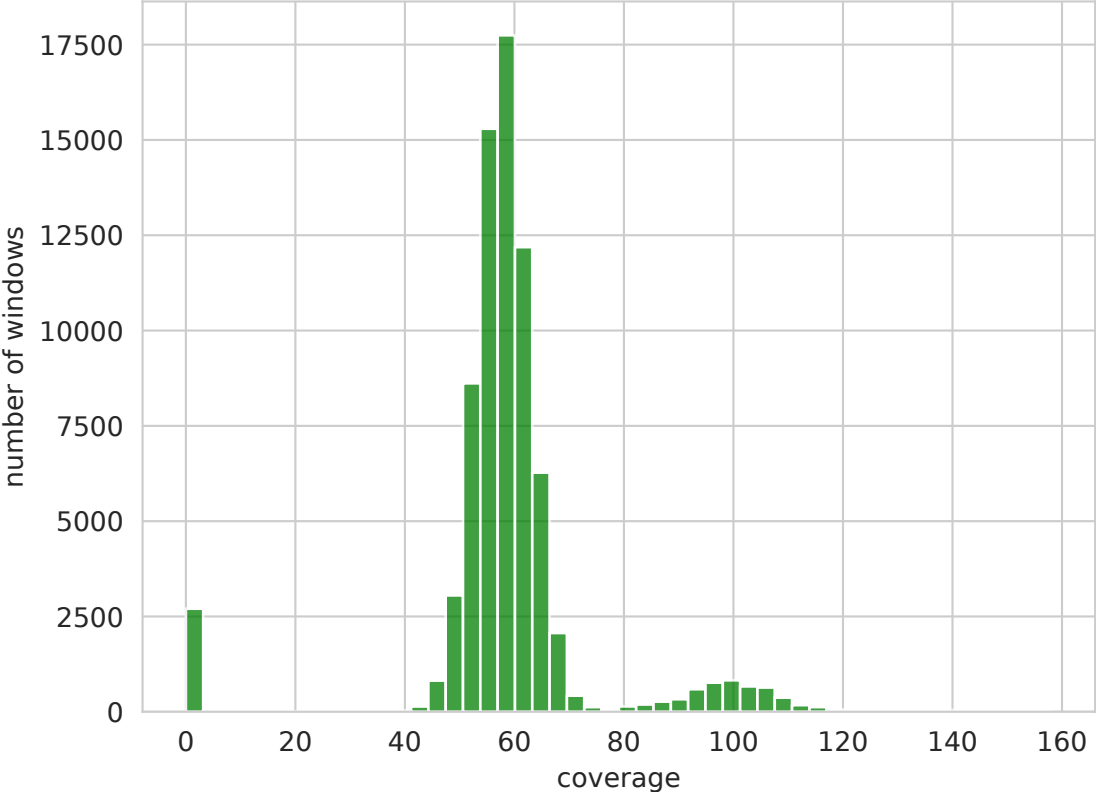
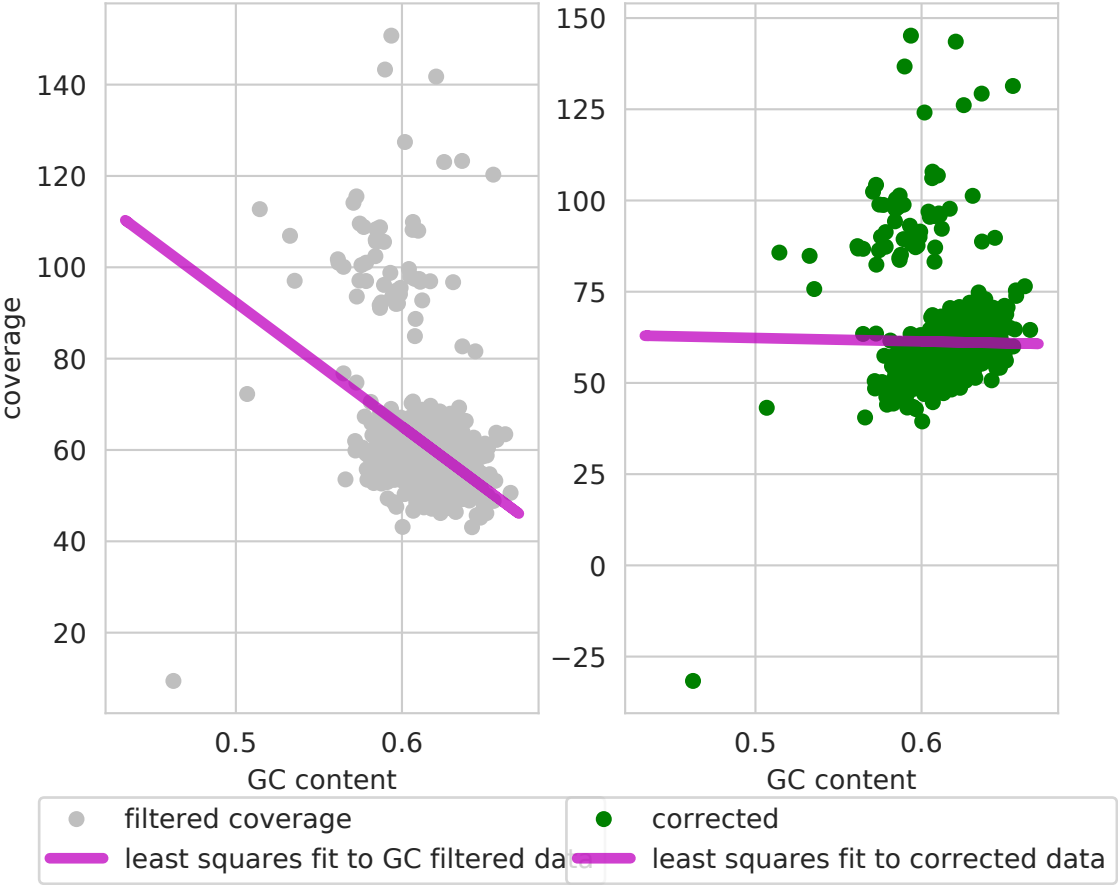


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0724/GCF_005938745.1_ASM593874v1_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0724-A
iRep: n/a r^2 : 0.85 avg. cov: 59.11 % windows: 96.38



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-A
iRep: n/a raw iRep: 1.42 GC r²: 0.21 GC bias: -57.18 avg. cov: 59.11 % 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-A
iRep: n/a un-filtered iRep: 1.42 r^2 : 0.85 avg. cov: 59.11 % windows: 96.38

