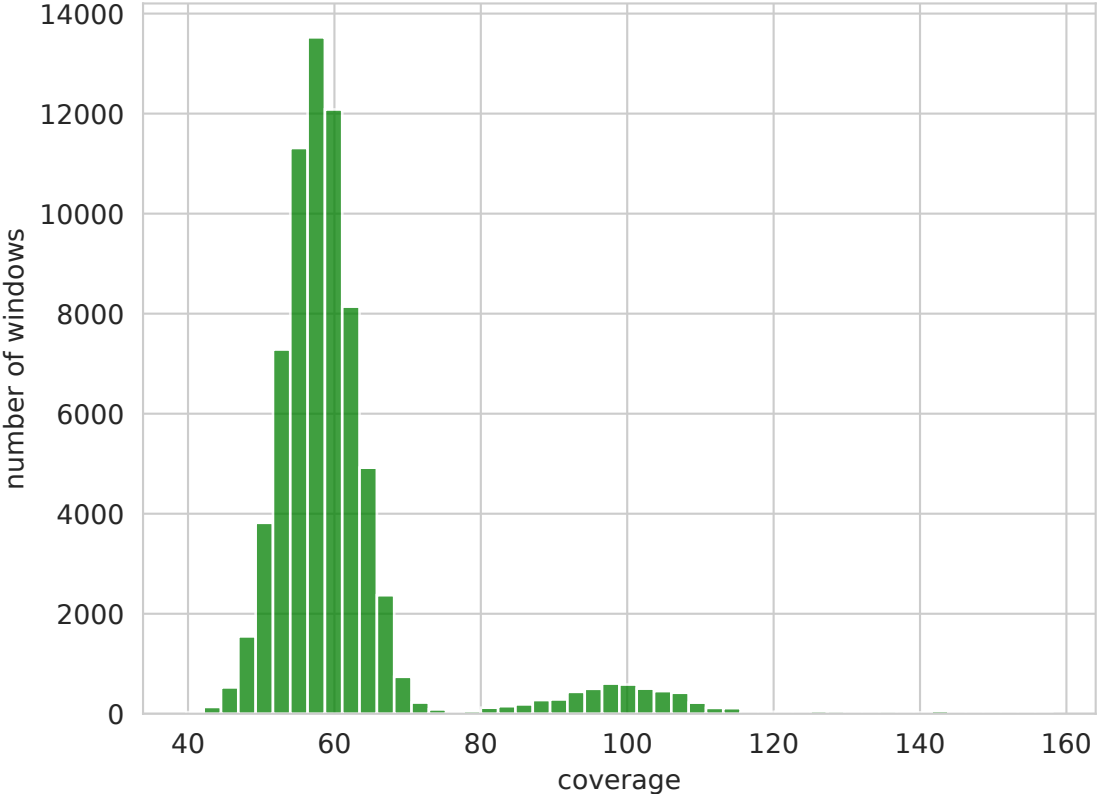
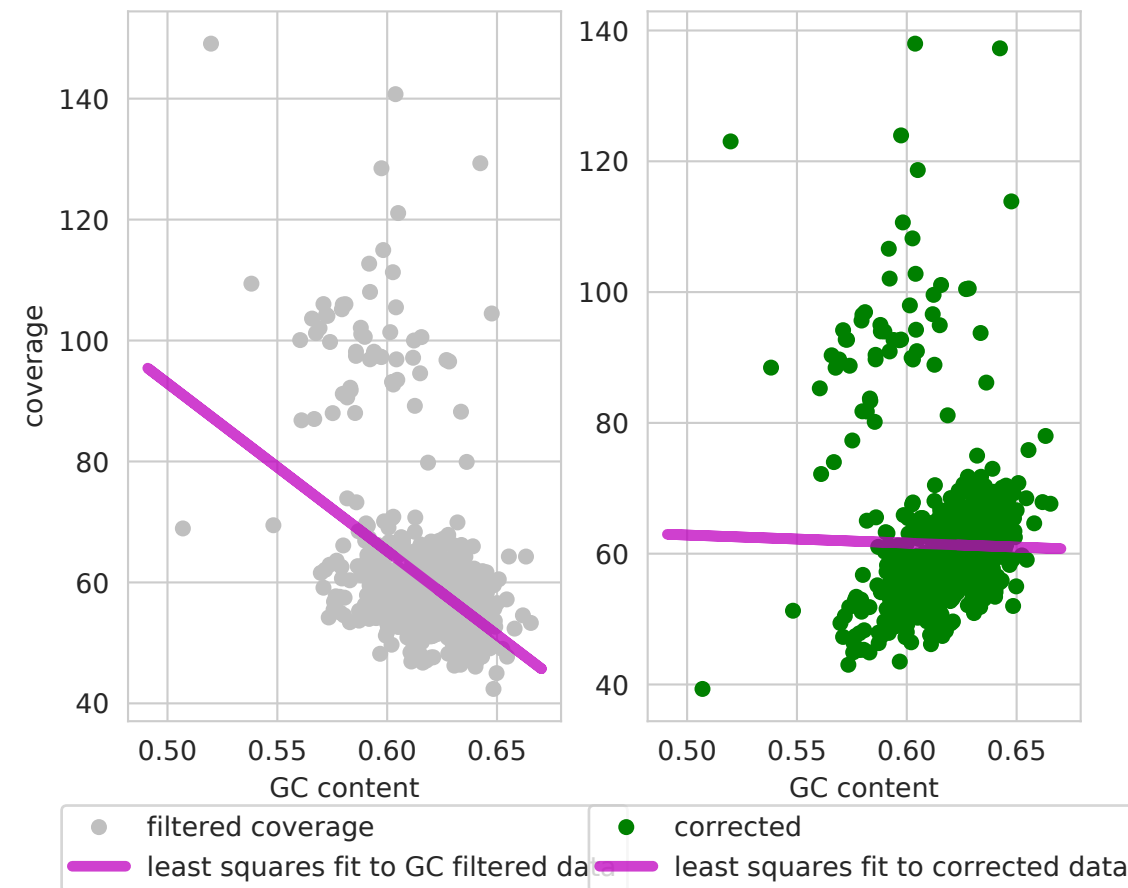


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0724/GCF_005938745_2_ASM593874v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0724-A
iRep: n/a r^2 : 0.86 avg. cov: 61.41 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0724/GCF_005938745_2_ASM593874v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0724-A
iRep: n/a raw iRep: 1.41 GC r^2 : 0.22 GC bias: -61.66 avg. cov: 61.41 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0724/GCF_005938745_2_ASM593874v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0724-A
iRep: n/a un-filtered iRep: 1.42 r^2: 0.86 avg. cov: 61.41 % windows: 100.00

