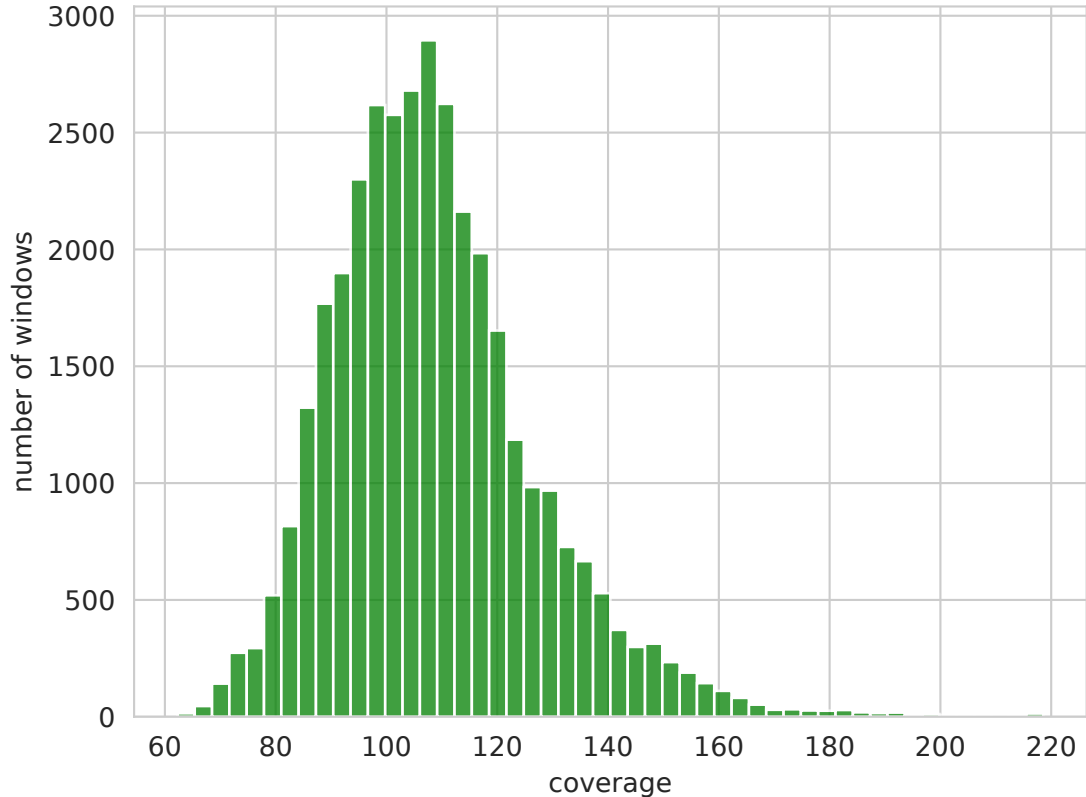
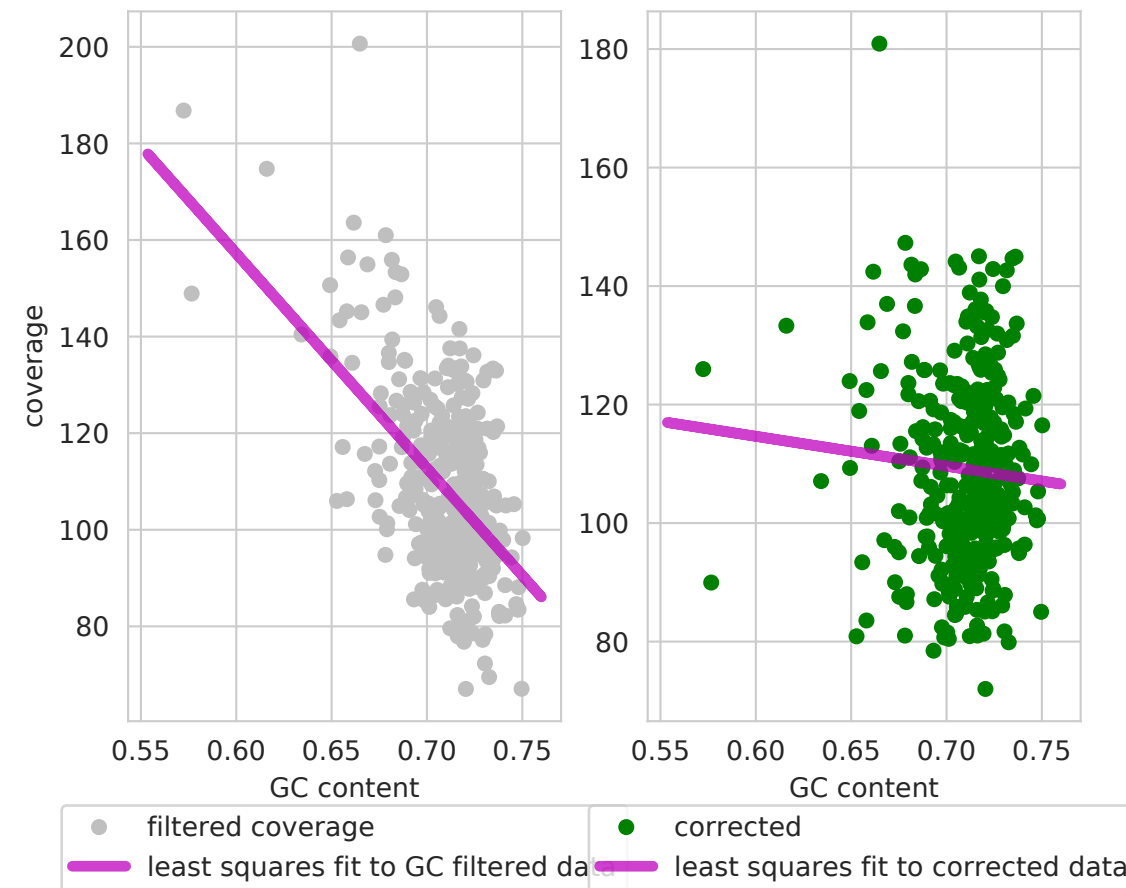


genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0715/GCF_005938005_2_ASM593800v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0715-D
iRep: 1.54 r^2 : 0.99 avg. cov: 108.64 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0715/GCF_005938005_2_ASM593800v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0715-D
iRep: 1.54 raw iRep: 1.62 GC r^2 : 0.30 GC bias: -134.49 avg. cov: 108.64 % windows: 100.00



genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0715/GCF_005938005_2_ASM593800v2_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0715-D
iRep: 1.54 un-filtered iRep: 1.54 r^2: 0.99 avg. cov: 108.64 % windows: 100.00

