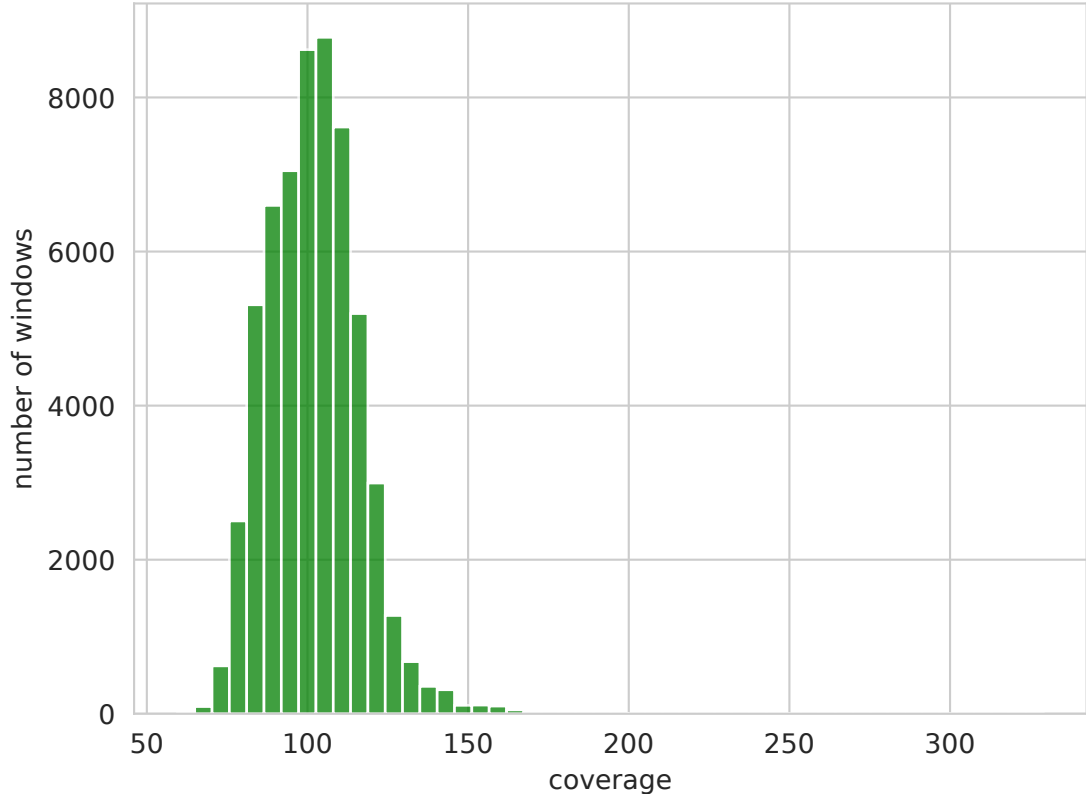genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065_3_ASM200706v3_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-D

iRep: 1.48   r^2: 0.99   avg. cov: 102.34   % windows: 100.00
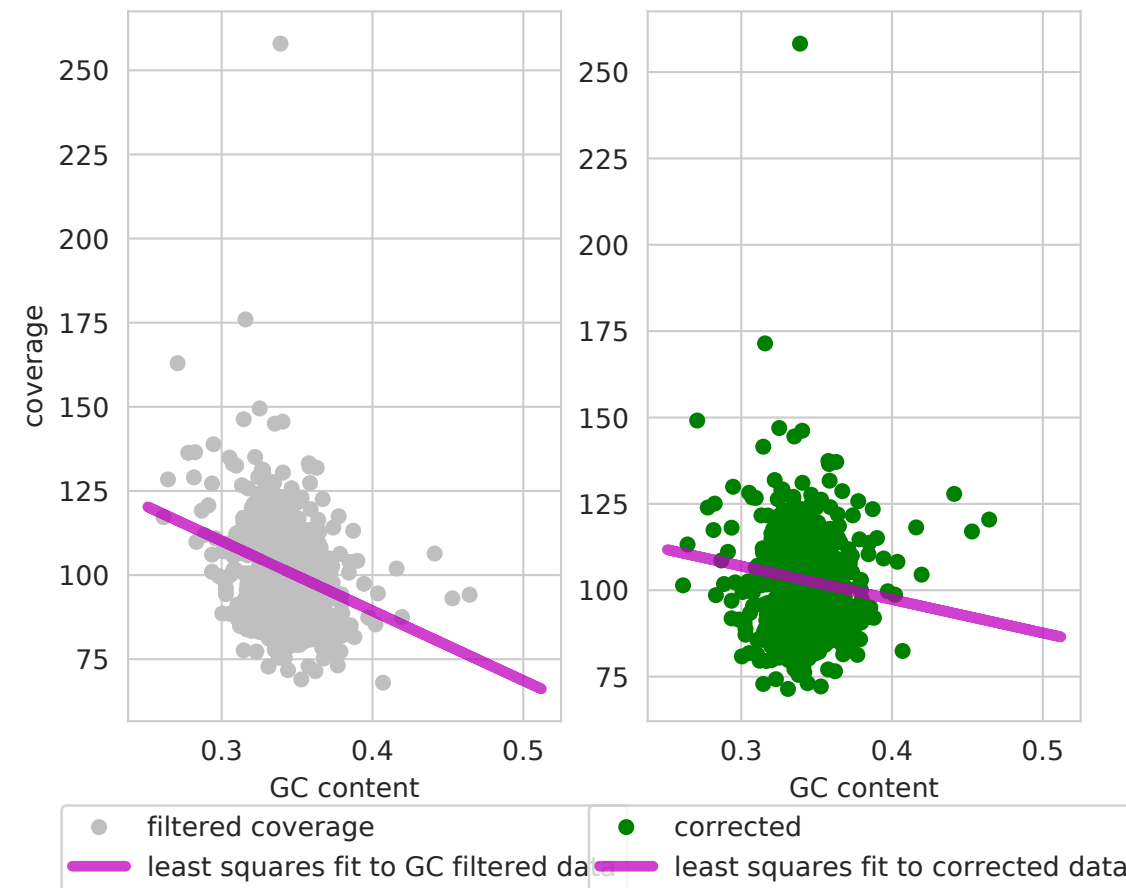
genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065_3_ASM200706v3_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-D

iRep: 1.48   raw iRep: 1.52   GC r^2: 0.12   GC bias: -24.70   avg. cov: 102.34   % windows: 100.00

genome: /N/dc2/projects/muri2/Task2/LTDE/data/genomes_ncbi/KBS0721/GCF_002007065_3_ASM200706v3_genomic sample: /N/dc2/projects/muri2/Task2/LTDE/data/bwa_bam/KBS0721-D
iRep: 1.48   un-filtered iRep: 1.48   r^2: 0.99   avg. cov: 102.34   % windows: 100.00