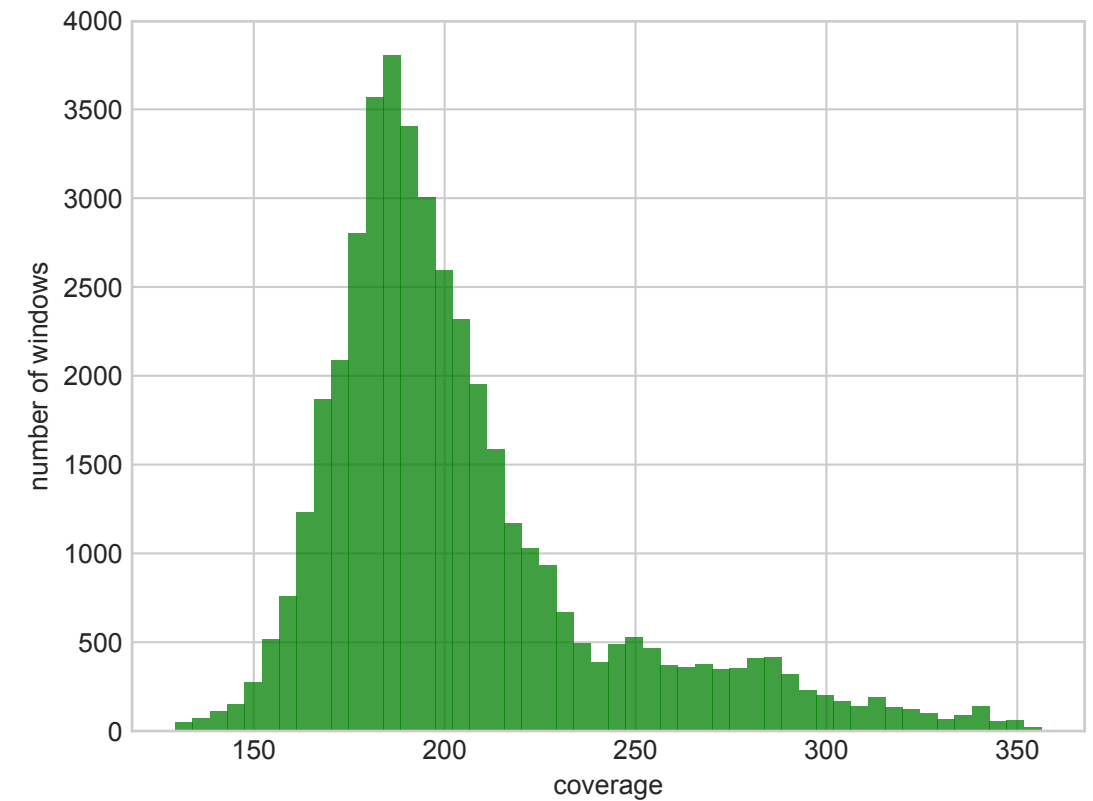
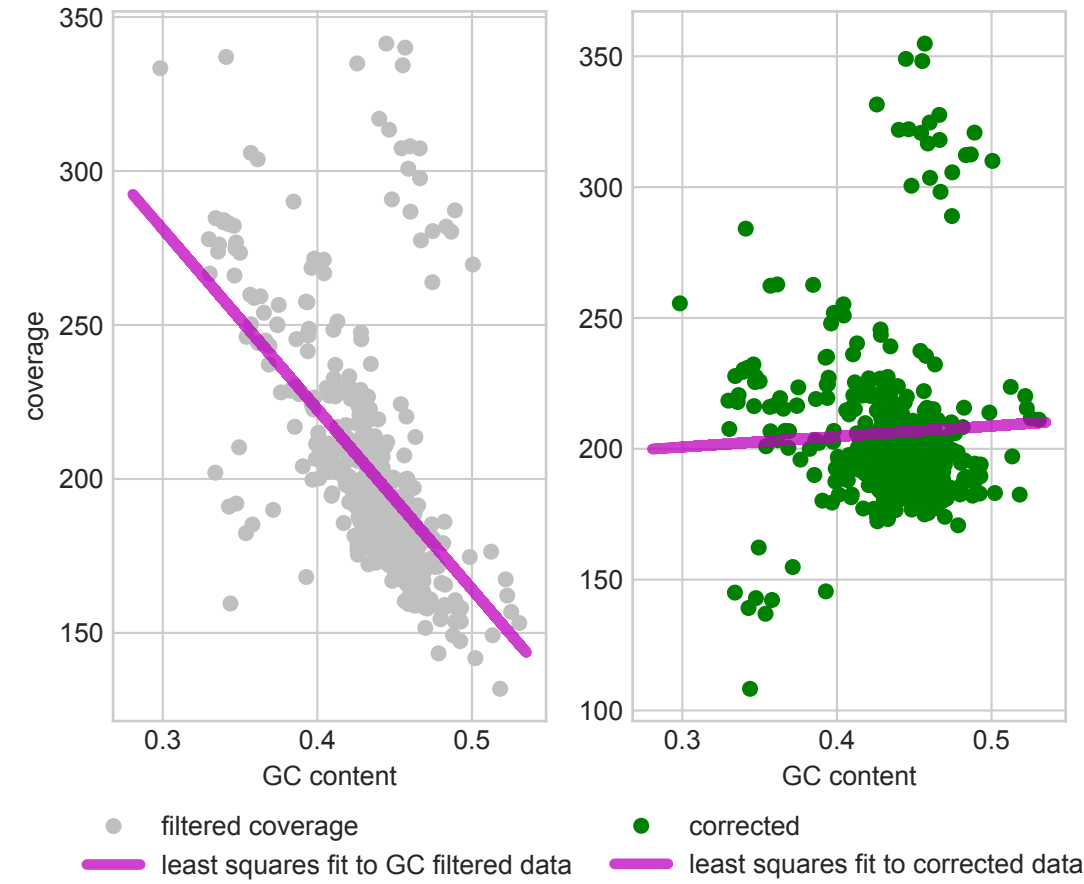


genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference\_assemblies\_task2/Bacillus\_subtilis\_NCIB\_3610/GCA\_002055965.1\_ASM205596v1\_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa\_fna/D300/Sample\_L2B5-300\_mapped\_sort\_NOdup\_sort\_merged\_sort  
iRep: 1.30 r^2: 0.90 avg. cov: 204.26 % windows: 100.00





genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference\_assemblies\_task2/Bacillus\_subtilis\_NCIB\_3610/GCA\_002055965.1\_ASM205596v1\_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa\_fna/D300/Sample\_L2B5-300\_mapped\_sort\_NOdup\_sort\_merged\_sort  
iRep: 1.30 un-filtered iRep: 1.30 r^2: 0.90 avg. cov: 204.26 % windows: 100.00

