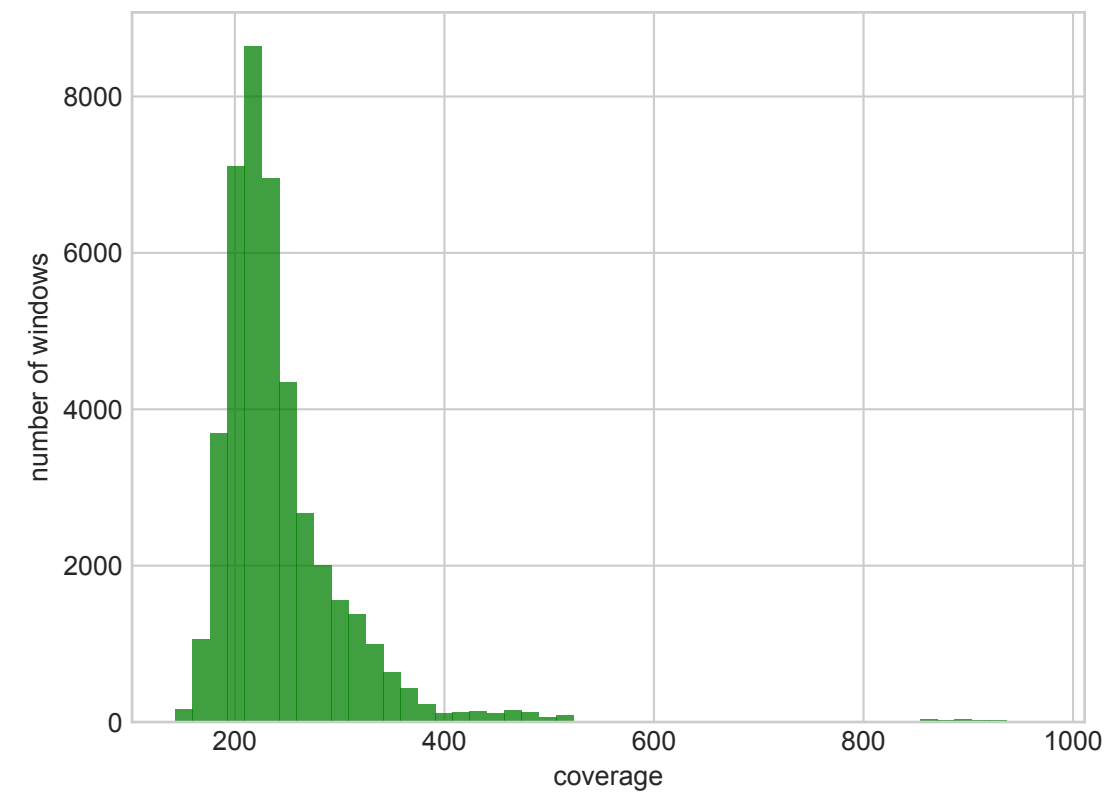
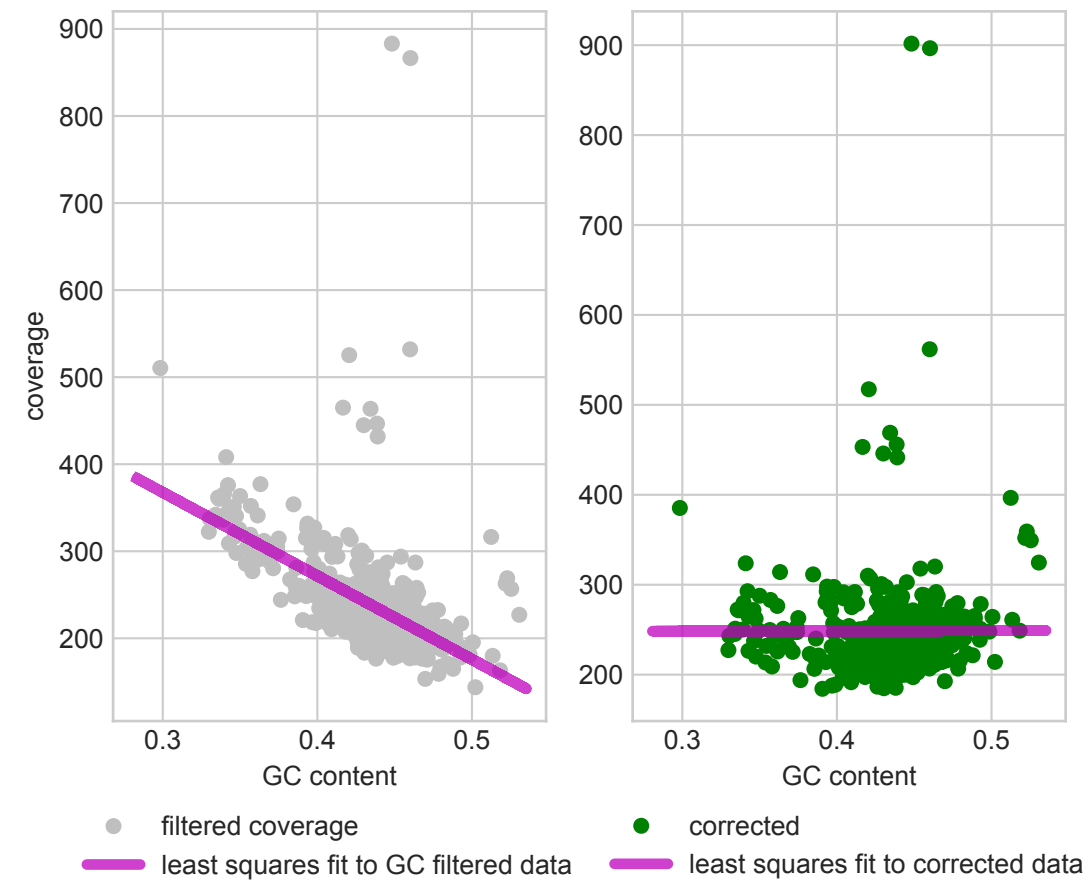


genome: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/reference\_assemblies\_task2/Bacillus\_subtilis\_NCIB\_3610/GCA\_002055965.1\_ASM205596v1\_genomic sample: /Users/WRShoemaker/GitHub/Task2/PoolPopSeq/data/bwa\_fna/D300/Sample\_L1B1-300\_mapped\_sort\_NOdup\_sort\_merged\_sort  
iRep: 1.43 r^2: 0.98 avg. cov: 244.10 % 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\_L1B1-300\_mapped\_sort\_NOdup\_sort\_merged\_sort  
iRep: 1.43 un-filtered iRep: 1.43 r<sup>2</sup>: 0.98 avg. cov: 244.10 % windows: 100.00

