genome: /Users/WRShoemaker/GitHub/Bacillus_Evol_Timeseries/data/Bacillus_subtilis_NCIB_3610/GCA_002055965.1_ASM205596v1_genomic sample: /Users/WRShoemaker/GitHub/Bacillus_Evol_Timeseries/data/bwa_sam_merged/Sample_L0S2-100

ptr: 2.06   avg. cov: 113.53

- coverage
- consensus Ori: 4,242,480
- consensus Ter: 1,942,720
- cumulative GC skew

genome: /Users/WRShoemaker/GitHub/Bacillus_Evol_Timeseries/data/Bacillus_subtilis_NCIB_3610/GCA_002055965.1_ASM205596v1_genomic sample: /Users/WRShoemaker/GitHub/Bacillus_Evol_Timeseries/data/bwa_sam_merged/Sample_L0S2-100

ptr: 2.06   avg. cov: 113.53

- coverage
- consensus Ori: 4,242,480
- consensus Ter: 1,942,720
- median filter
- cumulative GC skew