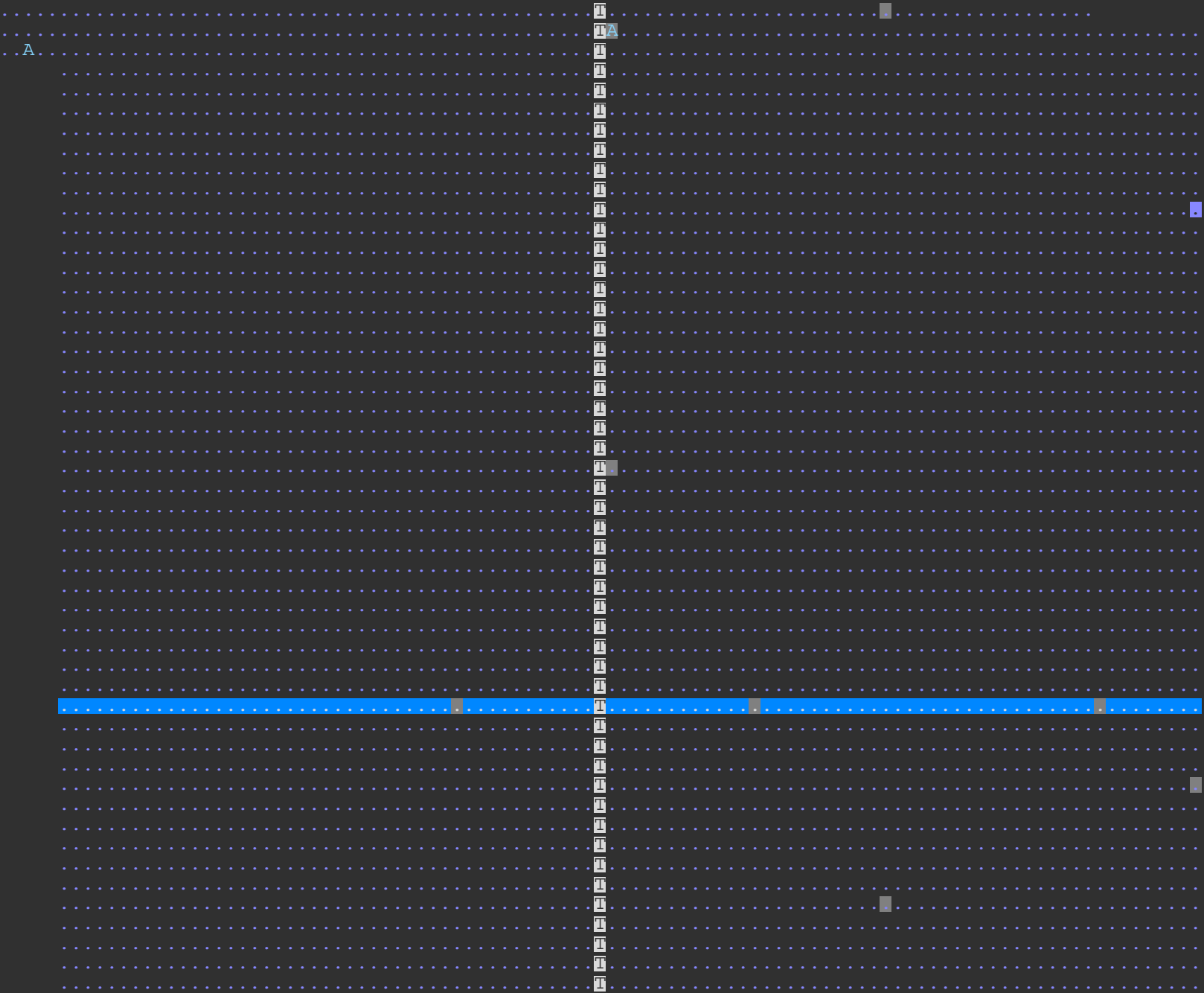


*-----6k-----12k-----18k-----24k-----3

SRR13500958.ivar_trim.sorted.bam@2; Reads: 910



SRR13500958.vcf.gz#3; N: 1

SRR13500958

nCoV-2019.primer.bed#4; N: 1

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GCA_009858895.3_ASM985889v3_genomic.200409.gff#5; N: 2

id-MN908947.3:1..265

MN908947.3:1..29903

CTT TA ATCT TTCTCTAAAC AACTTTAAATCT T T CT TCACTC CT CAT CTTA T CACTCAC CA TATAATTAATAACTAATTACT T

50 60 70 80 90 100 110 120 130 140 1

0 .10 .20 .30 .40 .50 .60 .70 .80 .90 1

MN908947.3:50-150; 101 bp; 1.0 bp/char

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