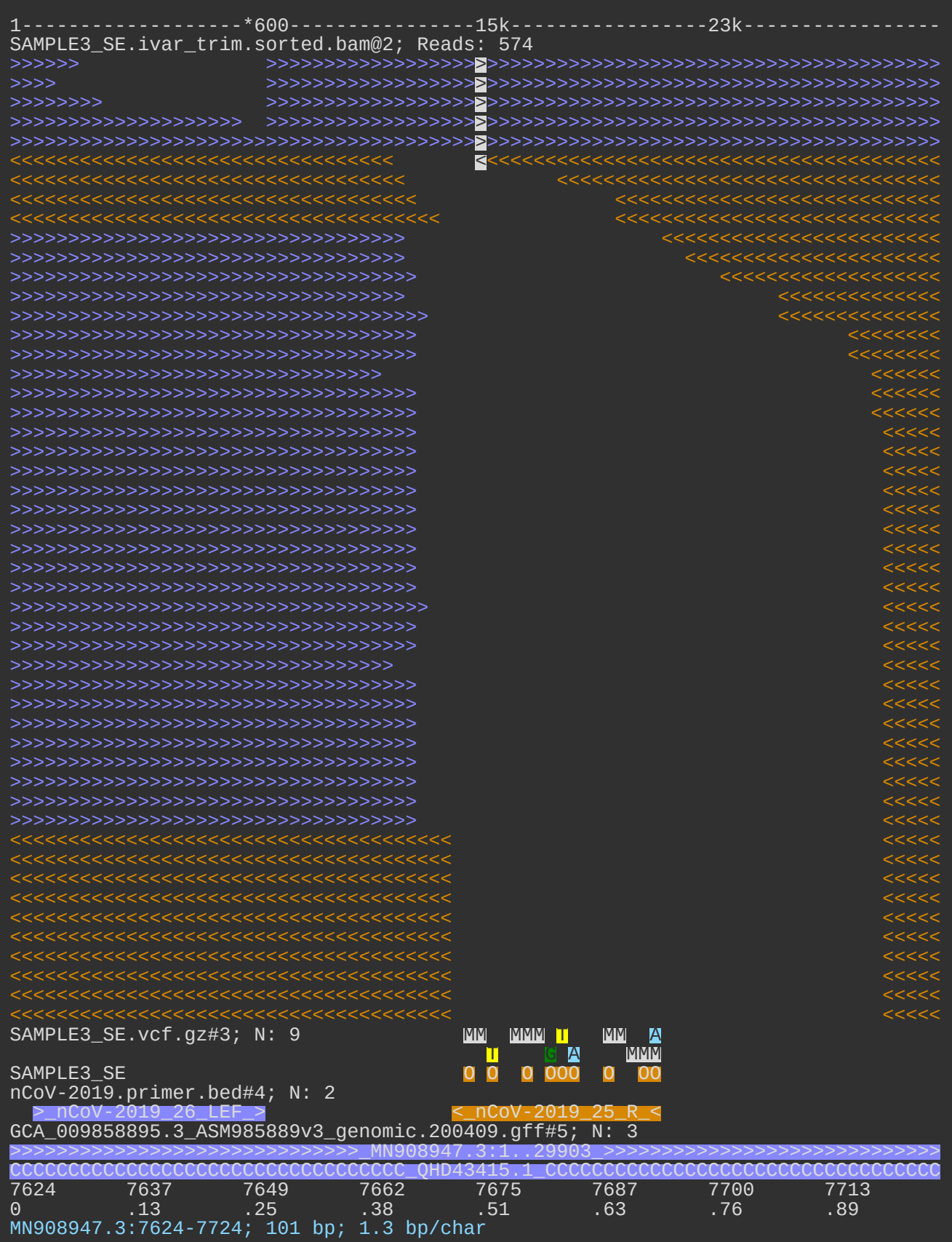
1----------*600----------15k----------23k----------
SAMPLE3_SE.ivar_trim.sorted.bam@2; Reads: 574
SAMPLE3_SE.vcf.gz#3; N: 9
SAMPLE3_SE
nCoV-2019.primer.bed#4; N: 2
>_nCoV-2019_26_LEF_>
<_nCoV-2019_25_R_<
GCA_009858895.3_ASM985889v3_genomic.200409.gff#5; N: 3
>>>