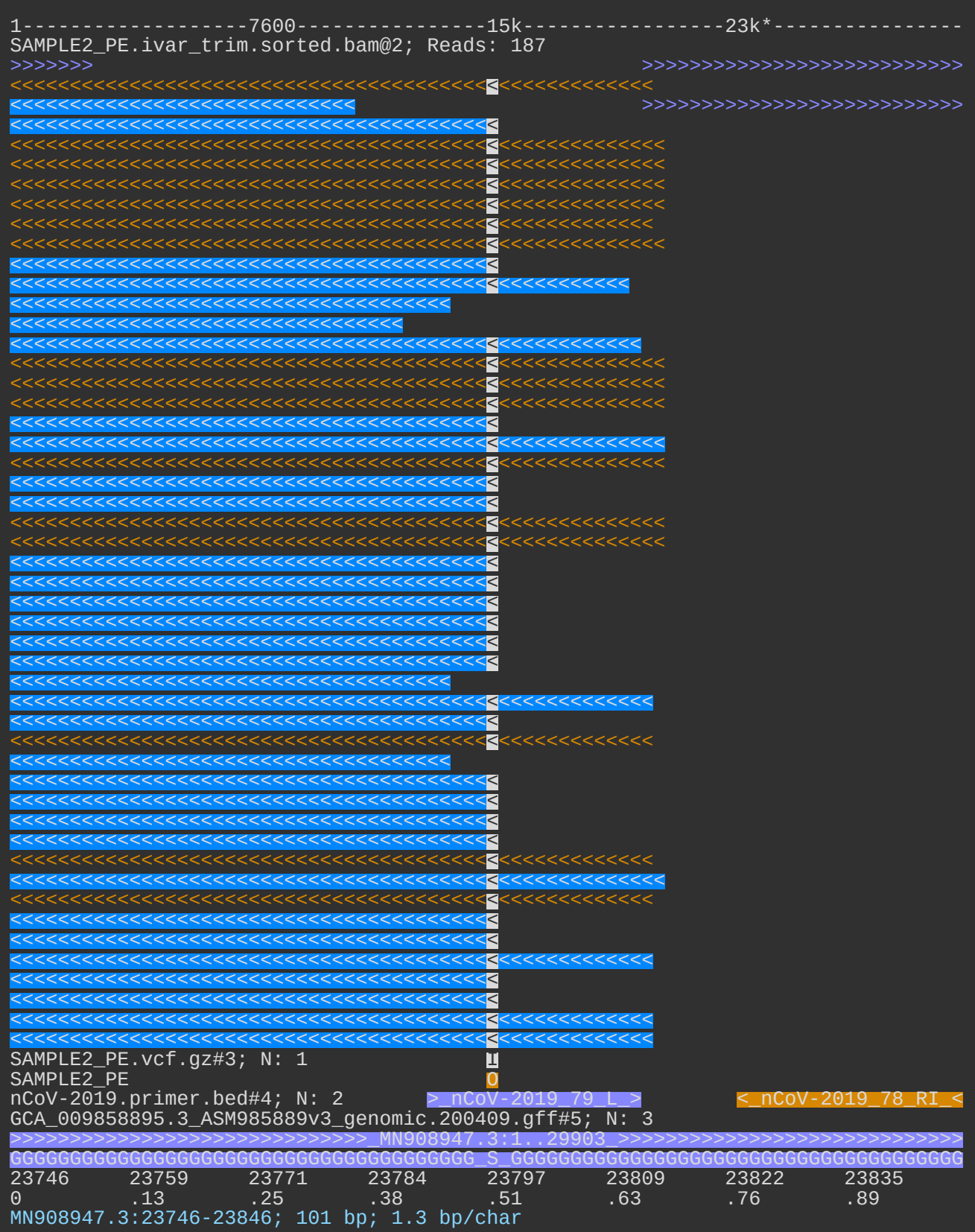
1----------7600----------15k----------23k*----------
SAMPLE2_PE.ivar_trim.sorted.bam@2; Reads: 187
SAMPLE2_PE.vcf.gz#3; N: 1
SAMPLE2_PE
nCoV-2019.primer.bed#4; N: 2
> nCoV-2019_79_L >
< nCoV-2019_78_RI_<
GCA_009858895.3_ASM985889v3_genomic.200409.gff#5; N: 3
>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>MN908947.3:1..29903_>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>
GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG_S_GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG
23746 23759 23771 23784 23797 23809 23822 23835
0 .13 .25 .38 .51 .63 .76 .89
MN908947.3:23746-23846; 101 bp; 1.3 bp/char