

1*-----6k-----12k-----18k-----24k-----3

SRR13500958.ivar_trim.sorted.bam@2; Reads: 1804

SRR13500958.vcf.gz#3; N: 1

SRR13500958

nCoV-2019.primer.bed#4; N: 0

GCA_009858895.3_ASM985889v3_genomic.200409.gff#5; N: 4

id-MN908947.3:1..265

MN908947.3:1..29903

CCCCCCC QHD43415.1 CCCCCC

GGGGGGGGG orflab GGGGGGGGG

TCTTACGTTTCCTCCCTTTTCACCCATCATCAACACATCTAATTTCTCCGCTTACC AAAATAAATATAACCTTCTCCCTTTTCAA

191 201 211 221 231 241 251 261 271 281 2

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

MN908947.3:191-291; 101 bp; 1.0 bp/char