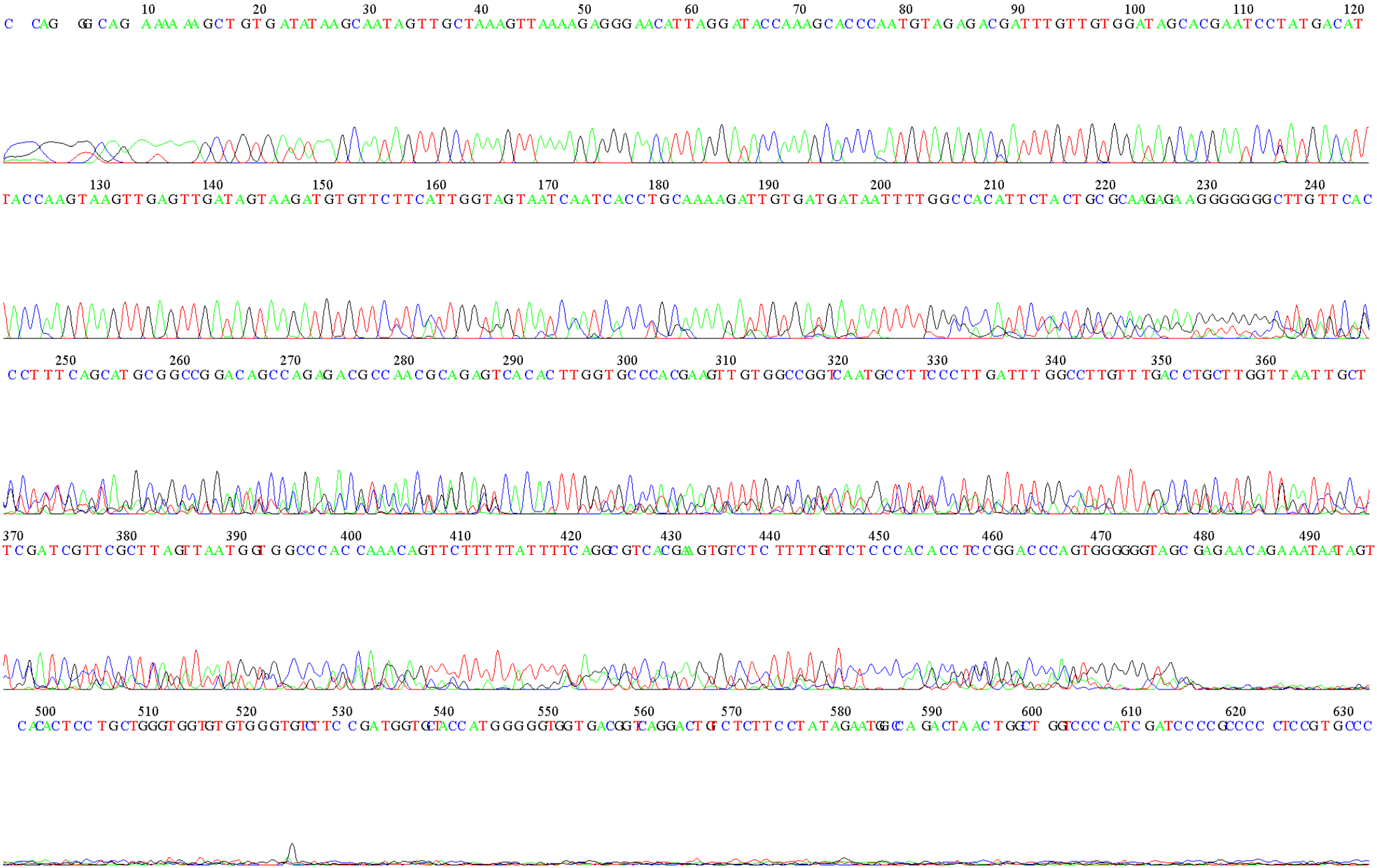
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C CAG GG CAG AAAA AA GCT GT G ATAT AA GC AAT AG TT GC TAAAGTT AAAAG AGGG AACATT AGG AT ACC AAAGC AC CC AAT GT AG AG AC GATTT GTT GT GG AT AGC AC GAAT CC CT AT GAC AT
130 140 150 160 170 180 190 200 210 220 230 240
TAC CAAGT AAGTT G AGTT GAT AGT AAGAT GT GTT C TTC ATT GGT AGT AAT CAAT CAC CT GC AAAAG ATT GT GAT GAT AATTTT GGC CAC ATT CTA CT GC GC AAGAGA AG GGGGGGC TTG TT CAC
250 260 270 280 290 300 310 320 330 340 350 360
C CT TT CAGCAT GC G GCCG GAC AGCC AG AG AC G CC AA C G CA G AGT CA C AC TT GGT GC CCA CGAAGTT GT GGCCGGTCAATGC CT TC CC TT GATTT GGCCTTGTT TGAC CTGCTT GGTT AATT GCT
370 380 390 400 410 420 430 440 450 460 470 480 490
T C GA T C GTT C GCT T AGT T AAT G GT G GCCC AC CAAACAGTTCTTTTTATTTTCAGGC GTCACGAA GTGTCTC TTTTGTTCTCCCACACC TCCGGACCCAGTGGGGGGTAGC GAGAACAGAAATAATAGT
500 510 520 530 540 550 560 570 580 590 600 610 620 630
CACACT CC TGCTGGGTGGTGTGTGGGTGTCTTC C GA TGGTGCTACC AT GGGGGTGGT GACGGTCAGGACT GC TCTCTT C CT AT AGAATGGCCA GACTAAC T GGCT GGTCCC CATC GATCCCCGCCCC CTCCGT GCCC

File: C67_78_Rev.ab1 *Run Ended: 2020/10/22 19:42:50* *Signal G:362 A:861 C:504 T:895*

Sample: C67_78_Rev *Lane: 41* *Base spacing: 14.745782* *1981 bases in 21104 scans* *Page 2 of 2*