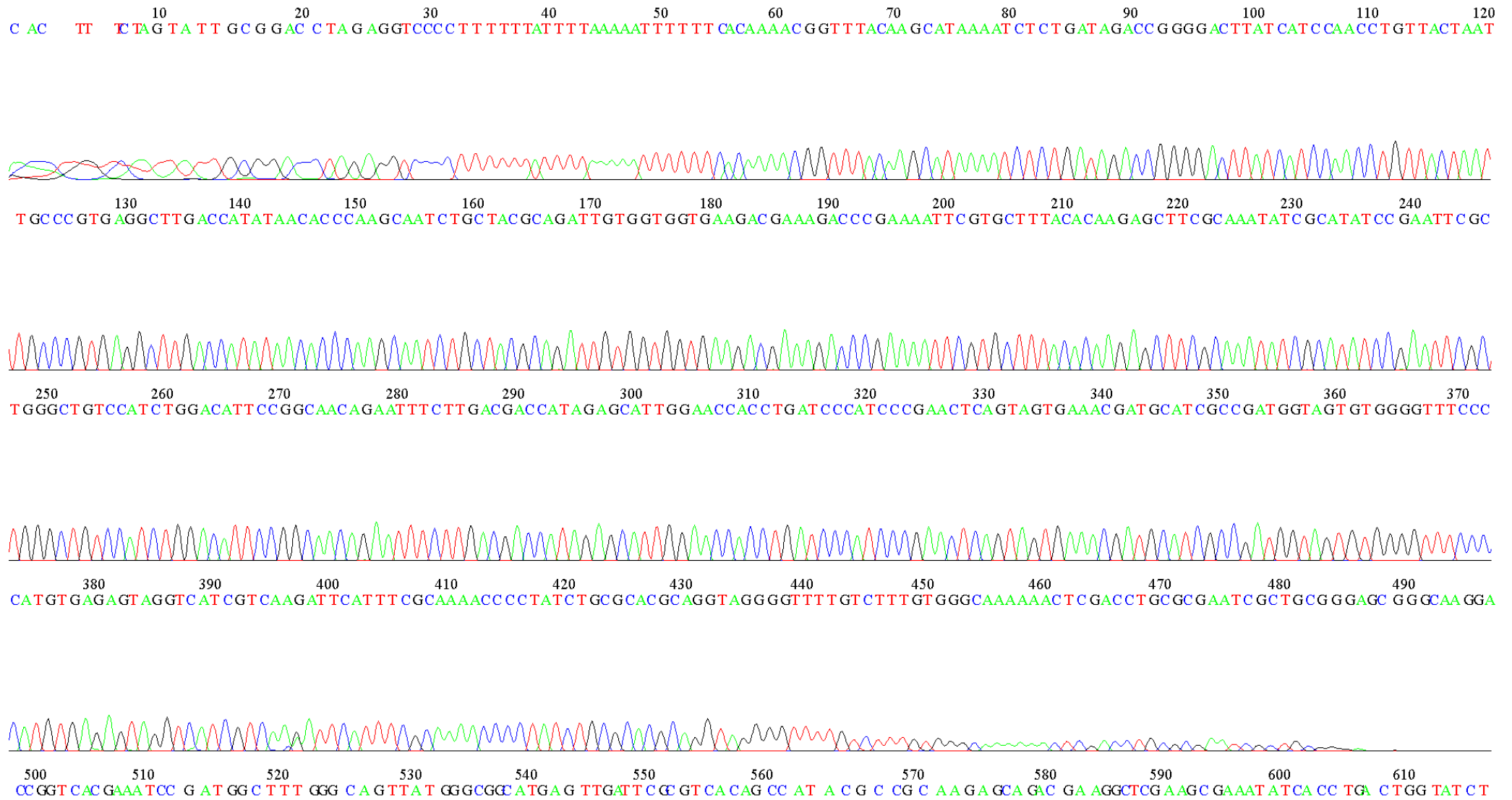


File: 22CCRAA000_E01_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:1335 A:2284 C:3021 T:2545
Sample: 22CCRAA000_E01_premix Lane: 7 Base spacing: 15.180631 1197 bases in 16301 scans Page 1 of 2



620 630 640 650 660 670 680 690 700 710 720 730
AGGA CG TCCACA GCA A CT AC CCAG TGGAC GTACT AG CGATGAA GTGTAGC AG AAGC CGGCC CCGAC GCGGGGGGC TGGAG G CAG ACTAAA CTAG CTC G AG ATTAGC A GGGGCTACA TTGC T

740 750 760 770 780 790 800 810 820 830 840 850 860
GCC G TAGGCTGG CGGCTG TA CATGCCGAC GA GAATGATCAA TAGCA GGTGAC GATGACCG CCCAG CAGC CAGGGT G AAGTG ATC GGG AGC CATTA TTA GAC GGGGAC GC AA AG CGAAC TGG CC

870 880 890 900 910 920 930 940 950 960 970 980 990
AGG CGATTGA TTTCCTC GGGCC GGGGTCGTGTTTGTATT GAAAA CG CGAAAC G AGTAT GACG TA GGAGACGAT TCCTTAG CG GAACT TGC GCC A CGCTTTT TAAGT G CT C G G G AAA C TTTT

1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120
AGTTACATC CGT AGAGGAGA GGT TTGCAAC CTGTAAACATTTT ACC A ATCTTG GAGAGGCCATAT GTTCTCC AAC ATGTA A GGCT G AGAGGG TTCCTT TGGG CCGGGG AA TGTG C C T

1130 1140 1150 1160 1170 1180 1190
TG TG TG TCGAGGGG C G GTCTTA A A G AC CTTCCTGC C AA AAGT CAGAA T TCTTT T GAC C