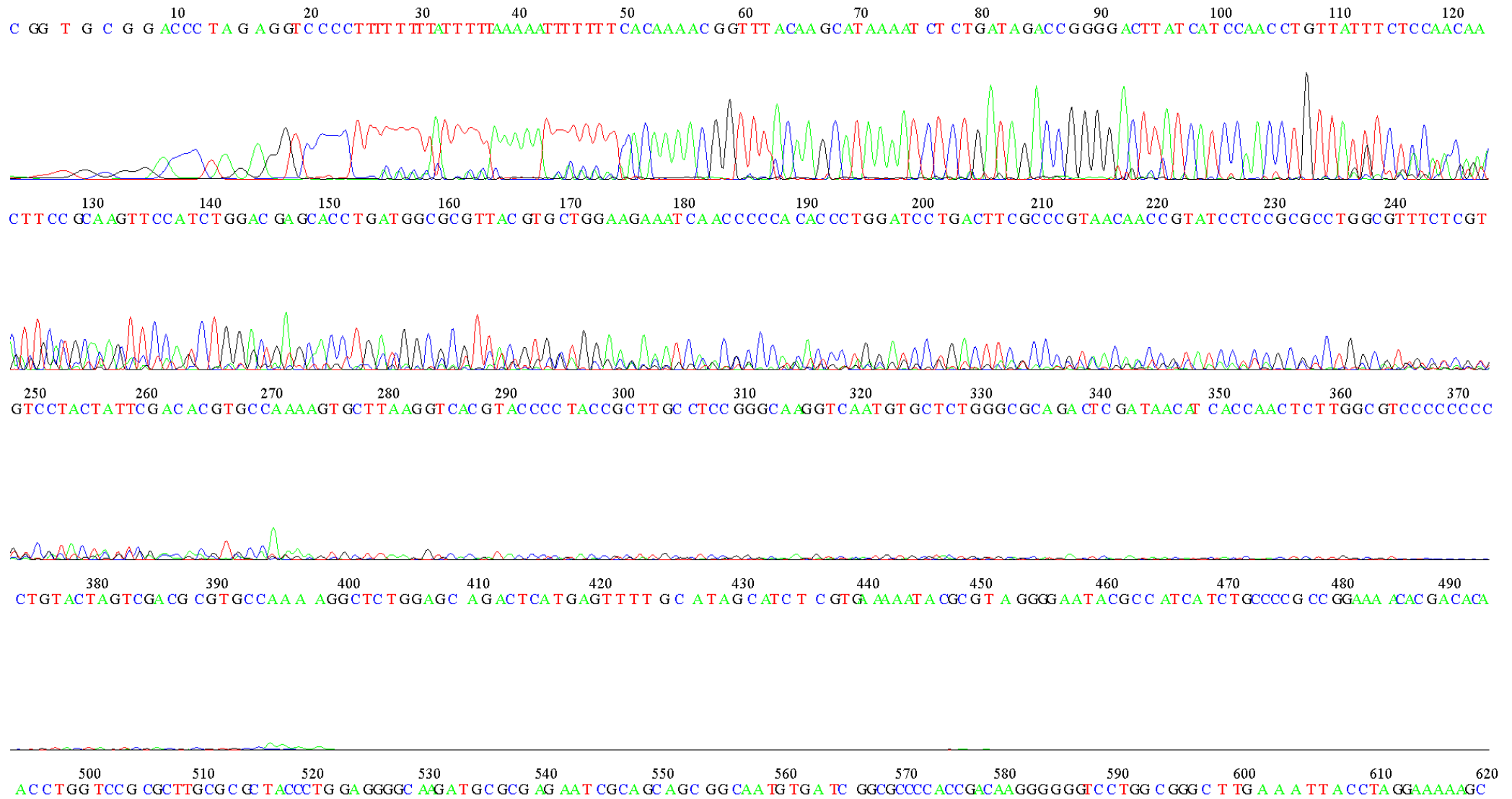


File: 22CCRAA000_H04_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:1555 A:2745 C:4175 T:3461
Sample: 22CCRAA000_H04_premix Lane: 18 Base spacing: 15.215738 841 bases in 16300 scans Page 1 of 2



630 640 650 660 670 680 690 700 710 720 730 740
TGAACGCCCAAGAA GAACCGA CCCATGCGCCGGGAGCACGTAA T GATACAGGA AACTTG TGCTCGA CCTGGTTGGATTAG GAGGA CAA GGG ACTG CCGAGTGCGCC GATACCTAGATT C GGA

750 760 770 780 790 800 810 820 830 840
TGATTTTGGT ATATTGCCCCCCCCCCTAGTAAG AAAAAA C GGGCC TGT TATTGT AAG AG AACTCGATCGA CACAG GGAACGGATG GT ATA
