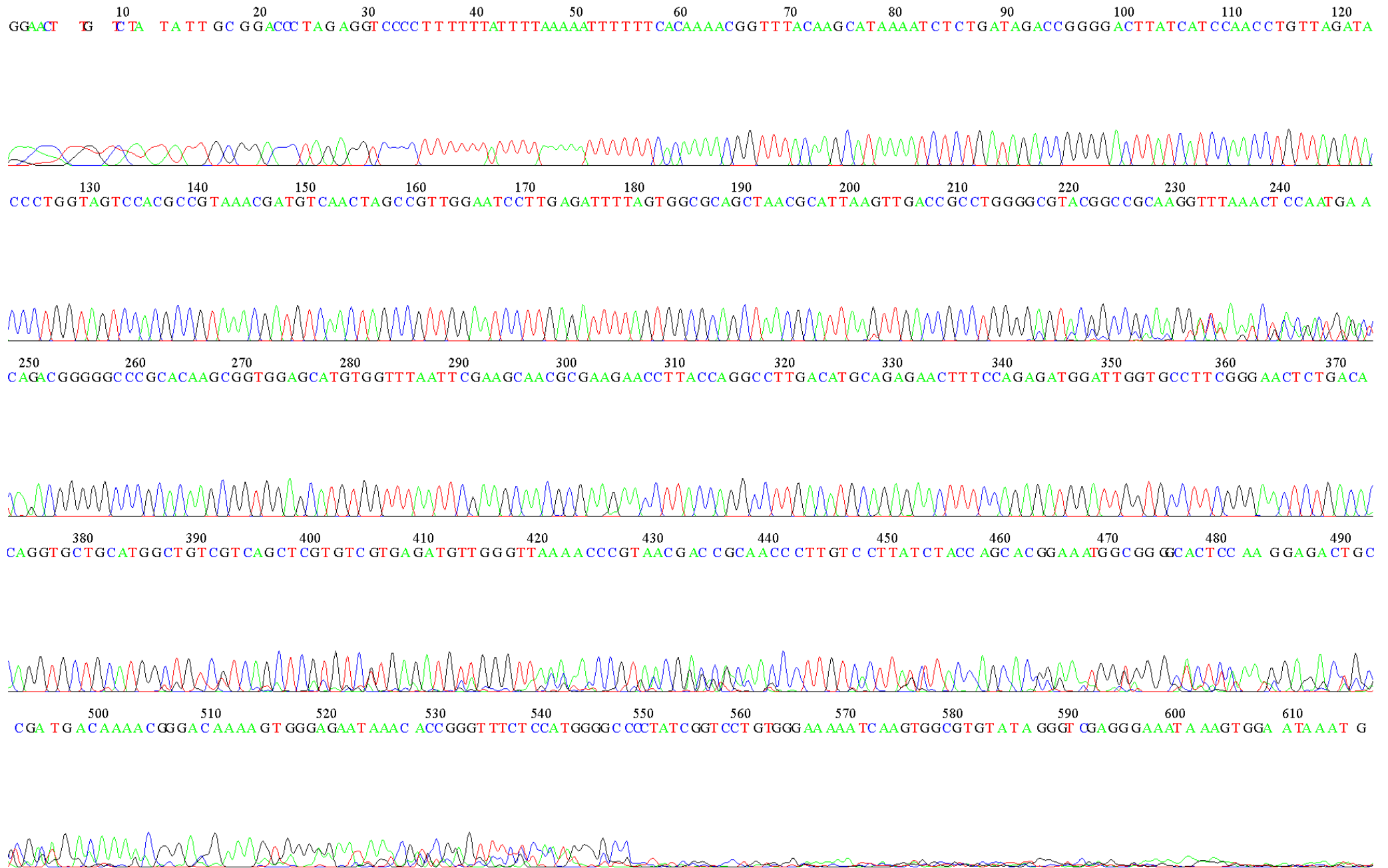


File: 22CCRAA000_G05_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:1377 A:1857 C:2274 T:1984
Sample: 22CCRAA000_G05_premix Lane: 35 Base spacing: 15.130663 2527 bases in 28104 scans Page 1 of 2



620 630 640 650 660 670 680 690 700 710 720 730
A CCGTAGCGT GA TAAAAA ACA GT CACAAAAACC GA AATAAT AC AACCCG GAAAC CCACAAACAT CGAA AA TACGAT G TAT C GGT G A G TGAAGTACAG AAG TCTAAT GTAAAGT GCC

740 750 760 770 780 790 800 810 820 830 840 850
TG C T TCAA CCCAA GGTATAGC AACTCAA ACCTGTAGGT TAGCACCGCCCG GGGATACATT GAATCAT AAGGAC A GTAGCTGAT C GGT C G CCGAGAC AA TA CG ACG AAAAAAGAA G C TTGATT G

860 870 880 890 900 910 920 930 940 950 960 970 980
TT ACTCTGC TGCTTTCAATTACTAAAAA CTTTGTGTG CACCTTCCT G AG TGC GAAAAAATAAGAGGA AAAAAA AAAGT AAAAAATAA A ACACACAT AAA GTAAC CGTAAAC AAA AAAAAA AAAAACTA TC

990 1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120 1130
TGCGCCACCGAA GAAAGTCTGCTATTAGGCTCTG TGGCATTCGGA GGGGTGACATGAAG AACAAACAT GATGGGCCATCCGACAATAACAGAGAA A ATCTTATCAACGGTC AATCAA TGA GTTTTGGGAGAC GA CGAGTATGTTG AA

1140 1150 1160 1170 1180 1190 1200 1210 1220 1230 1240 1250 1260 1270 1280
CAAAATCTCC CGCGTGT ACTAAAATAGAC CAAAC AAAAAA AACAAAAA C AAGA GGGAGGGG AAGAAATCTGTCTTT TTGAGTTAT TGTCTCCA A CGT AAAGAGGAGGTGCCCTCAT CCGTGTCCGTGCAAAAAA CTG