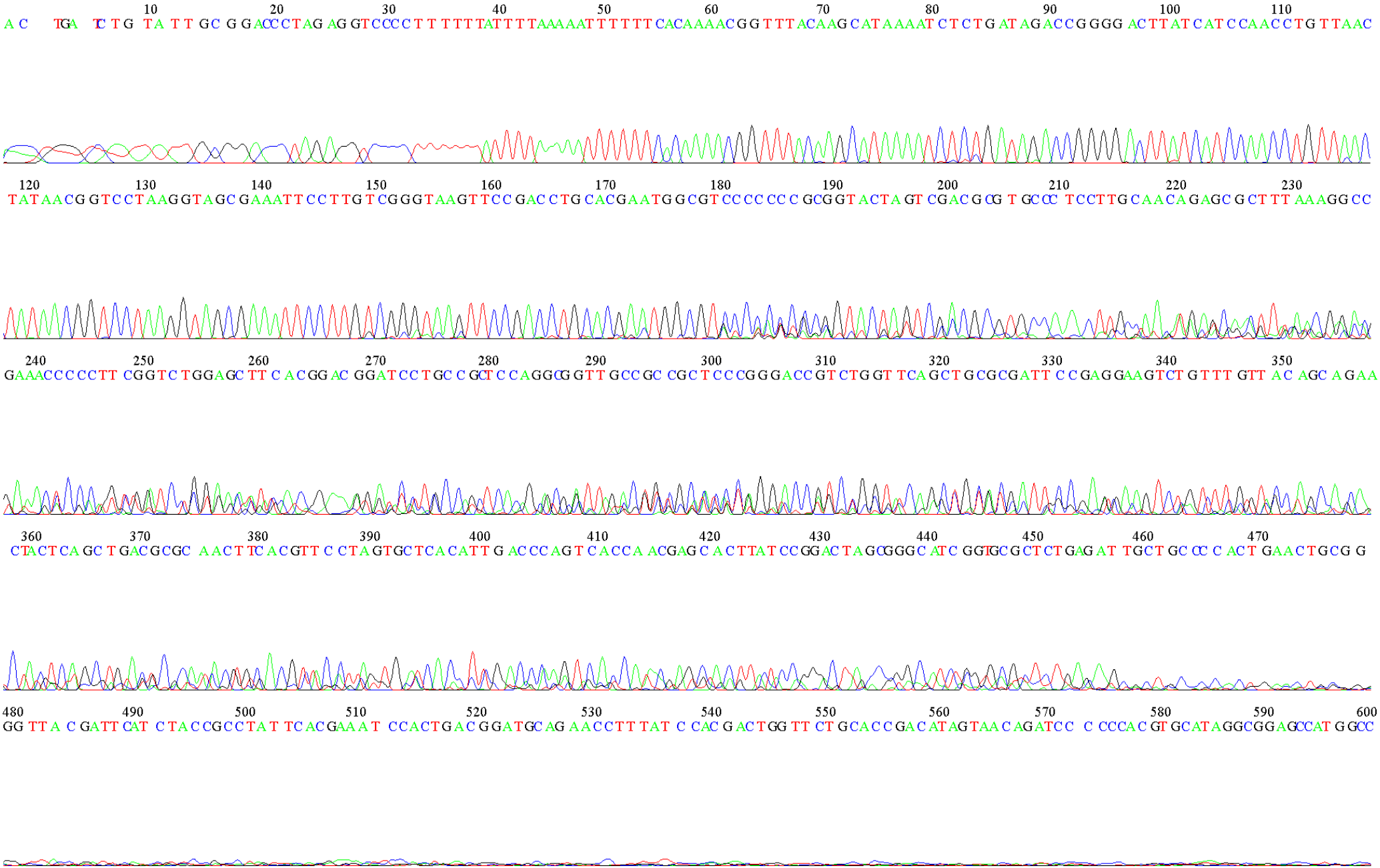
File: 22CCRAA000_A06_premix.ab1 *Run Ended: 2023/4/4 0:17:12* *Signal G:339 A:439 C:610 T:572*

Sample: 22CCRAA000_A06_premix *Lane: 48* *Base spacing: 15.168236* *1779 bases in 20949 scans* *Page 1 of 2*




610 620 630 640 650 660 670 680 690 700 710 720
GTCTA AT T AC A CC AA AT AATGGC G A CT ACCT A GC AT GG CC A T A T AT AAA A G A C A T C A A CTC GG ATCG G T GTCT ATT ATCT TTTTCC T CA CGTG G G TTG A A TTTT C CAAT T A C G TA TA GA GG T T
730 740 750 760 770 780 790 800 810 820 830 840
T G T C A G A A CAGT A C CCTTCTC T C T AC G T AG C C AG C AG GTGGCT GT CT AGG AC AT AGTGTT G GTTGT ACGGTG GA C T T A T T CT AT CA TT AC AC AG AT GAGT GAAGT CT GGCATATA GGTGAT
850 860 870 880 890 900 910 920 930 940 950 960 970
G AA C A GTC T TC CCCGG G C C G T A TT A CT CAA TC GC T A A TTC CC T GTTGA A ACATGTGGCC AATT A CT AACT TT G TGGT A A A GTTGC CGAGTCT GTT CTCCTAGTGT AGTTATG GTTCTAAC C CCT GT
980 990 1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100
T CT AAG GATGAGTCCT TA C T GGC TGTACT C GATGA A T AG AC TCAC ACTCTTT GT A CCCGAT TGG T A GCCG CTCTA A T GG TAT CAGCCGT CAGTT T CTTATGTGT AA TT GCGGGCT GTGACA GGTAT GCG T CC
1110 1120 1130 1140 1150 1160 1170 1180 1190 1200 1210 1220 1230 1240
C GC T AA CAGAA A C AAT ACTA CTCT A GTTA TACGAGT TAA GTCCATACATTGTGCGCC CGTC GGA GTATGATGCTTCTAT GC T GTT CA T ACGAT CGT TAGATGGCTCCGCAA AAAT ACCAAACCCAT T GTCATGGCGTAAT