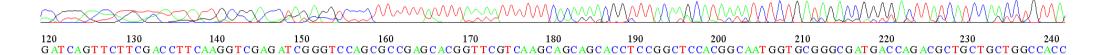
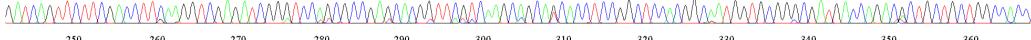
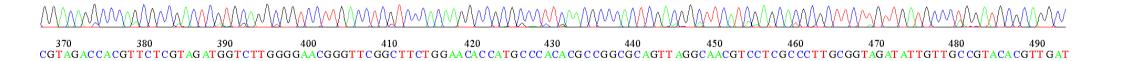


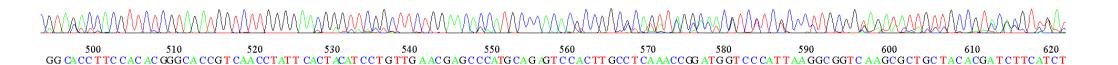
File: 22CCRAA000_G08_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:267 A:310 C:561 T:385

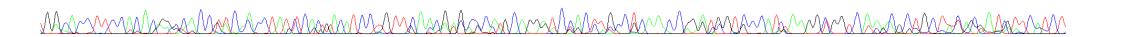




250 260 270 280 290 300 310 320 330 340 350 360
GG ACAG GC CCAGC GCCG ACTCGT GCAGG GCCTT GACCTCGT CCCACAAAGG CCGC GCCCTT CAA CGCCCACTC GACGG GCTT CAT C GAGCACGC GCTT C TT GTT GAT GC CCTG GAT GCCCAGGC







File: 22CCRAA000_G08_premix.ab1

Run Ended: 2023/4/4 0:17:12

Signal G:267 A:310 C:561 T:385

Sample: 22CCRAA000 G08 premix

Lane: 52

Base spacing: 15.351296

1315 bases in 16861 scans

Page 2 of 2

630 640 650 660 670 680 690 700 710 720 730 740 GCACAA C GAACAAC G CAT G

870 880 890 900 910 920 930 940 950 960 97

980 990 1000 1010 1020 1030 1040 1050 1060 1070

TGGGTAG AGGGGT GG ATT AAATT ATT CT GAG AA CAT TGAT AC AGG T A TAA TA TT CG GAT GAGGC T GAG AT GGGC A CAGGGT T ATA AAAAA TAGG A AT CT AG AGT

1000 1100 1100 1100 1100

1080 1090 1100 1110 1120 1130 1140 1150 1160 1170 1180
AT A TA AT A AGA GT AGA ATA GT TCGT CAA AA AAGA GT TTT AAG GA GT G GG GG T CAA TCTG GC GGTT GAGG T G TTTA AA A AAAAA G ATT G GGTT TGTA TA C AG GAA AAG AAA