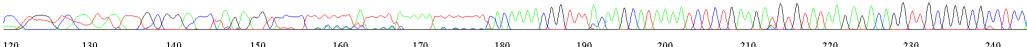
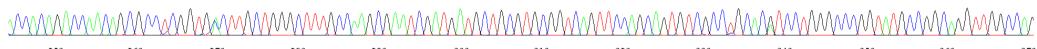


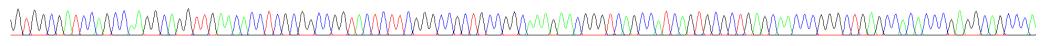
File: 22CCRAA000_H11_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:2571 A:2799 C:5271 T:3470

Sample: 22CCRAA000_H11_premix Lane: 81 Base spacing: 15.611872 1550 bases in 19126 scans Page 1 of 2

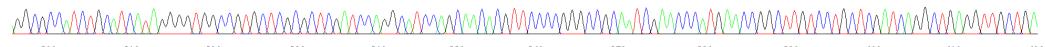




250 260 270 280 290 300 310 320 330 340 350 360 370 GTGGCGAT CCAGCCAAGGCAAGCCT CGCGGCCGGTACT CTTCGGGCCGCTGCCGGCAAAGAACGGGT CAGCCAT CAGT GCGCT GAGCAACCCGGGGCTGAGCCACA CCCGAGGCAGCCCCA



380 390 400 410 420 430 440 450 460 470 480 490
GGCGCCATCTGCATCATAGGGGTGGCCGTGCTCGCTCGATCCAGGCATCCAGCAACACGTTCCCCGGGCCGCAATCGAACCCATGAACCGGCTTGTCCTGCTCGATCAGGCTGAGGTTGCTGA



File: 22CCRAA000_H11_premix.ab1

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

Signal G:2571 A:2799 C:5271 T:3470

Sample: 22CCRAA000 H11 premix

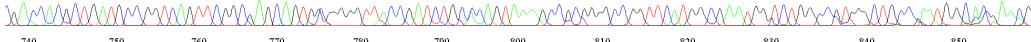
Lane: 81

Base spacing: 15.611872

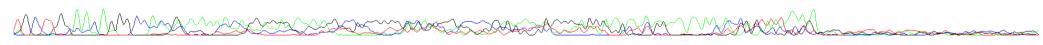
1550 bases in 19126 scans

Page 2 of 2

630 640 650 660 670 680 690 700 710 720 730
CGA CCACGCT GATGCC GGT CA GTTC GGCGAGCA GTGC GGGGTTACC GATCT GCACA GTAAA GCC GCGGGGCC GGTT CAT GG C GAATGG T CT GG CC GTGCC A AT G GC AC GAAC G









1080 1090 1100 1110 1120 1130 1140 1150 1160 1170 1180 1190 1200 G GGGTTAC CTAAA AT G C C CTCTCG TAGAAAGGG G G A AT A A A T T TTT TTT G T GCGGGAG GG GG GG GG GA AAAAA A TGA A CAT C CCC C GCC GA A C AGG GGGGGT TATAA A A CAT GC C G C C