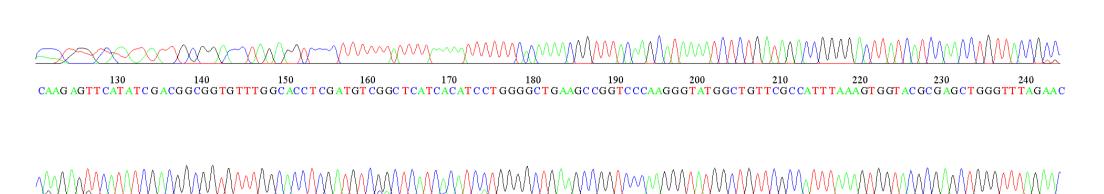
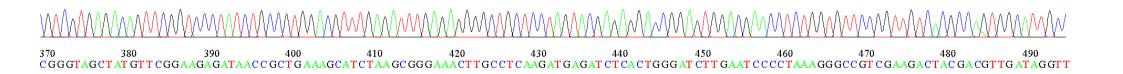
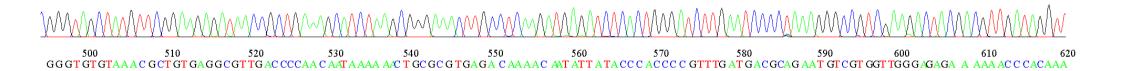


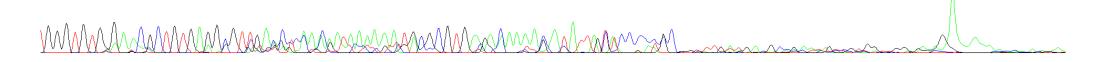
File: 22CCRAA000\_B05\_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:1128 A:1436 C:2238 T:1812











File: 22CCRAA000\_B05\_premix.ab1

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

Signal G:1128 A:1436 C:2238 T:1812

Sample: 22CCRAA000 B05 premix

*Lane: 45* 

Base spacing: 15.09988

1748 bases in 20247 scans

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630 640 650 660 670 680 690 700 710 720 730
ATA G G GTGT T CT C G A G A G C GCG AA TATAGTAG A T GAT A AGA A T C GT GTG G A TGC GC GG G A AGAT CT TCATAT A GCAA GGT GTT G G G A CGCC AAT A GAAC GGT A GAGAC CG C A G GTG

870 880 890 900 910 920 930 940 950 960 970 980
TCAGGACACC CGGA AAA GAA CATAACAA TGGGGGGGG AAT TACCCTT GCTA GAG GCCTT TTA G GGGTAAAA ATGCTGCCC AA GAGAGAGGCTC CCTA GGGGGGGGCC AGGTTGGC GT C CTAA AAGGG A

1130 1140 1150 1160 1170 1180 1190 1200 1210 1220 1230 1240 1250 1260 1270 1280 CCCCCTC C GACTT CT/GAAGAGTAT TTTA T AGAGAGAGGAGGAGGAGGAGGAGGAGGACCCC CC T TTTTGGTGAAACA GAGTGTTTT TTC ACACCCCCAT TCT TTT TGT G/GGGAC CC ATT ATTT CTT TGAATG T AA AAA AA AAGACCCA AC CCCCAAA