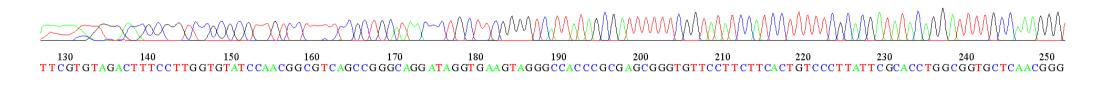
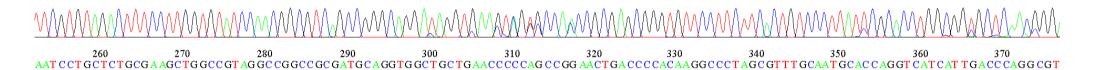
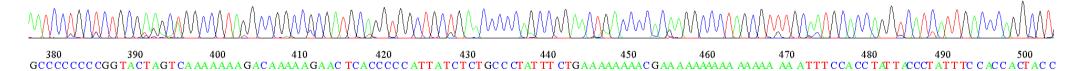


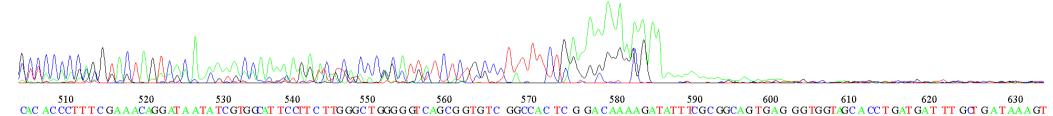
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10 20 30 40 50 60 70 80 90 100 110 120
TT A CT TGC T ACCG CG GT C CGC GC GTTTGTC CTT T TC CGC T GCATTAACCC T G CTTCCG GGGT CATTATAG CG AT TTTTT C GGTATAT C CAT C CTT T TT C GCACG AT AT ACA GG AT TTT GC CAAA GGG

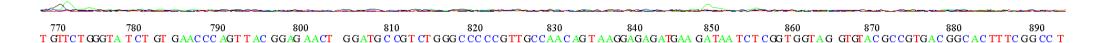




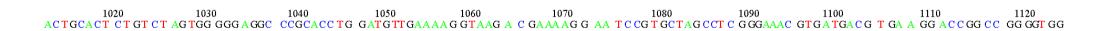




File: 22CCRAA000_A02_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:3044 A:2441 C:4790 T:4516 Sample: 22CCRAA000_A02_premix Base spacing: 15.270588 *Lane: 16* 1743 bases in 20425 scans AG GA A AAGTT GC A AG ACA AA GTATGAAG CE A COCG T C AC GGA C GGG CT GGCCATCCAGA CAT A CGAC AGA G ACAGA G C GA C GC CA C AAGA C AG GGTAACG GATCAGC G T C AC CG CT CA C CC GT GAT



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GC GAA A G C CGG GGG T T AGA CG GGGAAA CCTCA GAGA AA CAA TA AC AA A G C T C T CG GGG GC CTTG GTG GGGGT C T G TT A C C CGGC C T ACGGG GGAT C CACCT C ACCC G C G C G