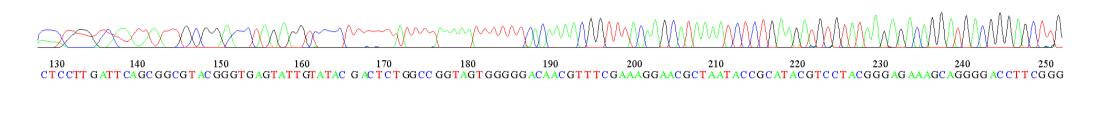
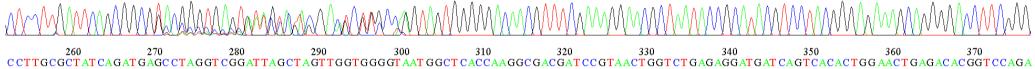
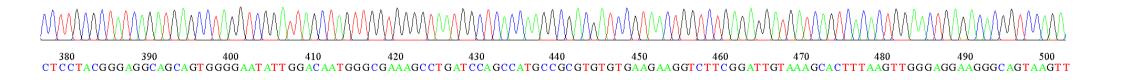


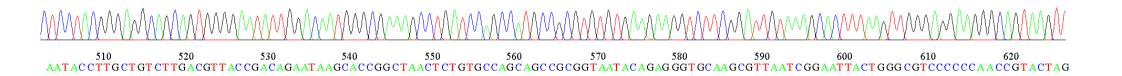
File: 22CCRAA000_B12_premix.ab1 Signal G:2541 A:4089 C:4833 T:4589 Run Ended: 2023/4/4 0:17:12

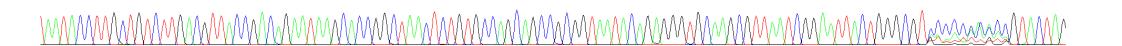
Sample: 22CCRAA000_B12_premix Base spacing: 15.42219 Page 1 of 2 Lane: 94 1552 bases in 17599 scans











File: 22CCRAA000_B12_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:2541 A:4089 C:4833 T:4589 Sample: 22CCRAA000_B12_premix Base spacing: 15.42219 Page 2 of 2 *Lane:* 94 1552 bases in 17599 scans

CAAA AAA C GAA AAGT TT GGAGTGTACGCA GCAC G C CCCTC G CT CTTTTC AGG C GAGCGG AGT AG C CGCAAG GA AGACT TCAA GGCCGGAGCGTGGTTCCAGC GCGCCGTACATA T AG AGGAAGCA CAAG GAAT AAGCACACGCGT AGCACACTA T AG AGGAAGCA CAAGA CAA