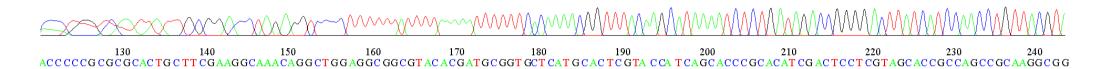
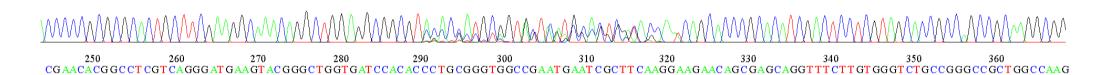
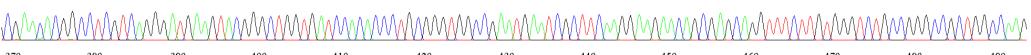


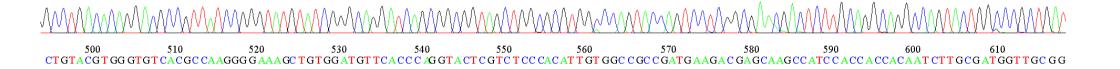
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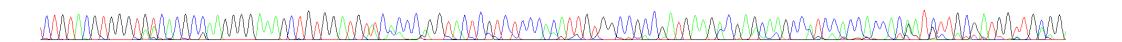






370 380 390 400 410 420 430 440 450 460 470 480 490 GCCTGACACAGGACGCCGTTATCCGGGTATGTATCGGGCAGGATCAGCGGGGTAGCTGGCCCGCTGAT





File: 22CCRAA000_C04_premix.ab1 Signal G:1399 A:1934 C:3248 T:2358 Run Ended: 2023/4/4 0:17:12 Sample: 22CCRAA000_C04_premix Base spacing: 14.995245 Page 2 of 2 *Lane*: 28 1575 bases in 19753 scans TT TT CCCAA C A AAAAAA A AGA A G GGGTC C CTC C CC CC CCCGGGC C CCA C GTAATGGTT G CA AAA AAAAAAC A AA AAAAAA AA TTCCC CG GGT G GG GC GGGT CG CCC CTT G AC CGG G GA AA

