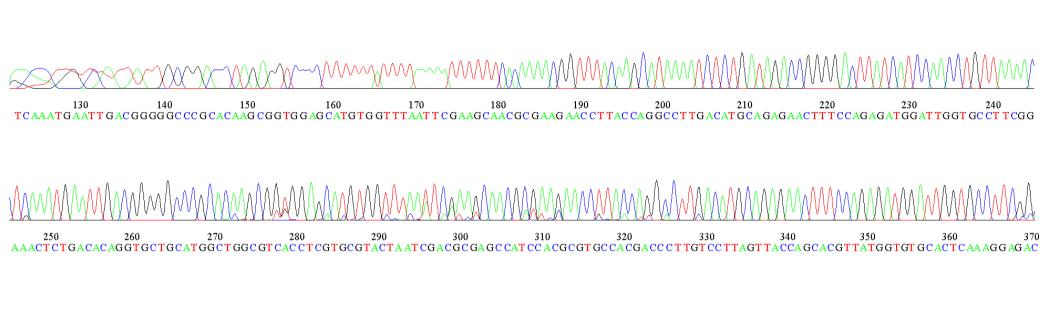
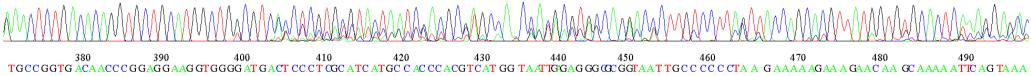
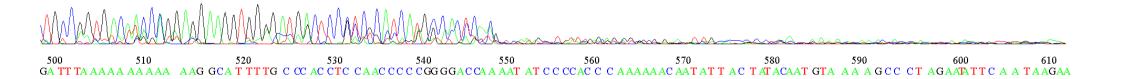


File: 22CCRAA000_A01_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:886 A:1178 C:1423 T:1115

Sample: 22CCRAA000_A01_premix Lane: 15 Base spacing: 15.215109 1077 bases in 13600 scans Page 1 of 2







Sample: 22CCRAA000_A01_premix Page 2 of 2 Base spacing: 15.215109 1077 bases in 13600 scans *Lane: 15* 740 750 760 770 780 790 800 810 820 830 840 850 CA CAAAG TA AAA AAAA AGTA G CCG AAC G TTGC AATGCT AT A AAA C A G AGA GG GC AGAT A ACTAAT GA AAA GG ACCAAG AGC C GC A AT ACA AAAAAA GACGT AG AA AA AAA C GA A ACTAGGAT GA AGGTGA AT CTT GA A AATT AAA A AAA AGAA AA AGAA AA AAATAC TACCCTGT AAATA TACA AAAA TGA GC AGG ATAAT GC CAACA C GT G GGA TTAA CTAA ATAA C CTCCAGGGA ATGAAATGG TCAACGT AAAA AC G GTAGA AAAAAAA GGG GT GGGCAG TACGGAA AAGGCAGT CATCTCCCCCCCT T C CAAA AAAAAA C TG GC

Signal G:886 A:1178 C:1423 T:1115

AA AA TAC

File: 22CCRAA000_A01_premix.ab1

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