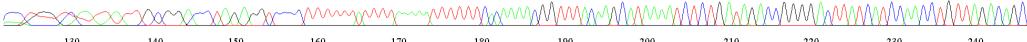
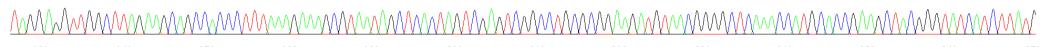


File: 22CCRAA000_A04_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:1054 A:1659 C:2159 T:2045

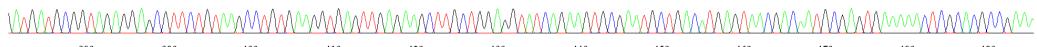
Sample: 22CCRAA000_A04_premix Lane: 32 Base spacing: 15.132 1396 bases in 16276 scans Page 1 of 2



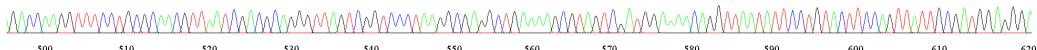
130 140 150 160 170 180 190 200 210 220 230 240
TAGGAGGTTGGCTTAGAAGCAGCCCCTTTAAAGAAAGCGTAATAGCTCACTAGTCGAGTCGGCTGCGCGGAAGATGTAACGGGGCTCAAACCATGCACCGAAGCTACGGGTATCATCTTATG

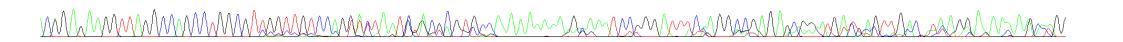


250 260 270 280 290 300 310 320 330 340 350 360 370
AT GAT GCGGT AGAGGAGC GTT CT GTAAGCCT GT GAAGGT GAGAAGCT TGC T GGAGGTAT CAGAAGT GC GAAT GCT GACAT GAGAAAACT C GCACGC C GAAA



380 390 400 410 420 430 440 450 460 470 480 490 GACCAAGGTTTCCT GCGCAACGTTAAT CGACGCAGGGTTAGT CGTCCCTAAGGCGAGGCT GAAAAGCGTAGT CGATGGAAAACAGGTTAAT AT TCCT GTACTT CCAGTTAT TGC GATGGAGGGA





File: 22CCRAA000_A04_premix.ab1

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

Signal G:1054 A:1659 C:2159 T:2045

Sample: 22CCRAA000_A04_premix

Lane: 32

650

Base spacing: 15.132

1396 bases in 16276 scans

Page 2 of 2

740

GTTGATA AG AAGC GACC A AC C A GA CT CC A AG AAT T TA A AA TGA G A GC GGG CC CT CC GCCGACC GAGGT AGTT AGGC AC GT GT AAATA C GC AG GT CAT AGAC A CA CTT GGA AGAC CG ACC GA

M

750 760 770 780 790 800 810 820 830 840 850 860

AG C AACAGAGAT CGAT AT CTAC OG GAGT AC GC GCG CCTGC AGA GCOCGAA A GACATAGAT ACG AAT C C C G T G CT TT A CGAG AAGG TAT GA C G CT C C G AC AGT AT ACT A GAGA G CAGAA GT GA GC AGA

870 880 890 900 910 920 930 940 950 960 970 980 990

GC GG ATT GT GG C TTACC GGGGGGGGGGGGGCCC GG C C T GAA C C G AA C GCT C A TT GT T GGA T G T GGGA GAA GA A C C C C C GG T C GG AA C ATT AA GGGCCG C GA AA C TT T A GGGA A C ATT GTT C C AA

1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120 1130 A TTT C GGGAAAAAAAAAA CCGC CGTTTATAAAGTG GGG G GTTTTT GTATTAAC CACAGA GAAAGC AA AAAAAATT GGGG TIT GTTCCCTTCCCGG GTTT CTT GCCA TTT TT AAAAA AAA A G AAAAA AAGGC GAC