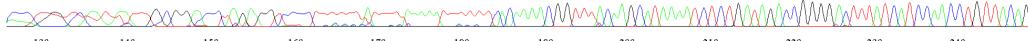
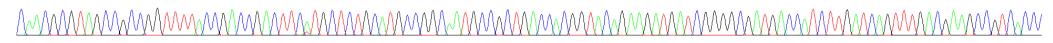


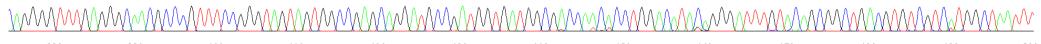
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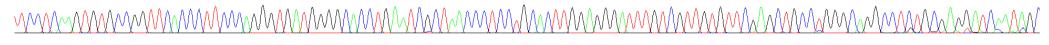
10 20 30 40 50 60 70 80 90 100 110 120 ACC TG TCTA T TATTTG C G GAAC C TA GAAG GFC C C CTTT TTT ATT TTAAA AAT IT TITTT (A CAAAAC G G T T T ACAA G C AT A AAAT CT C T G AT A G AC C G G G G ACTTT AT CAT C CAAC C T G T T ACA





250 260 270 280 290 300 310 320 330 340 350 360 370 CAGG GGTTTGAGGGCCAACGCCCGTTTCCG GGTAGTAGT GGCCGAGGAT GCCGATGGTGAT GGCCGACCACACCCT GGACAT CAGGG ATTTGTACAGCGGCT GACGT GT CGT CAT GGCTAATTT





510 520 530 540 550 560 570 580 590 600 610 620 AAGGGTGGTGCCAAATGCCCAAGGTCCGTATTTGCTGGGGATAAATGCCCAACGACGGCGATACTCCCGGGGACGGATTGCCGGCGATGGCGGAATTTGTA

File: 22CCRAA000_D04_premix.ab1

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

Signal G:2128 A:2824 C:4268 T:3064

Sample: 22CCRAA000_D04_premix

Lane: 26

Base spacing: 15.078618

1571 bases in 19759 scans

Page 2 of 2

630 640 650 660 670 680 690 700 710 720 730 740 C AA GG AA GG AA GG AA GAAT GG GG AA AT CCACCCTTT GG GG AT GC CG GG AG AT CT G GT CAT A C CCTT GAAA C GAG AA GT TT CAT GC GT AGAC AACCC GT T GG GGC GGAAAG GT T TT CTT GAA

