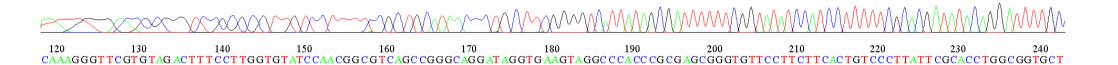
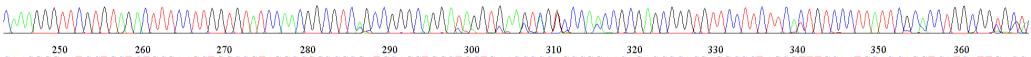


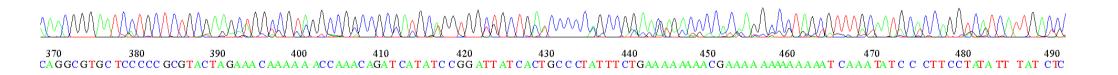
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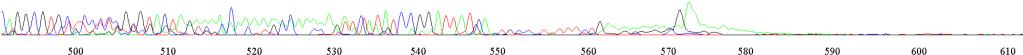
Sample: 22CCRAA000_A11_premix Lane: 95 Base spacing: -16.163063 1655 bases in 20860 scans Page 1 of 2





CA ACGGGÃAT C CT GCT CTGCGA AGCT GGCCGTAGGC CGCGAT GCAGGT GGCT GCTG AAC CC CCAGCCGGAACAGAC CC CA CGAGG CC CT AGCGTTTGCAAT GCAC CAGGT CATCATTGACC





500 510 520 530 540 550 560 570 580 590 600 610 A C AAC TT CTT TA C CAGAG AT C AACA GCCT ATA AAGGGCCC TT T GGAC AGT GG GGA GTGTA C G TG CT C GTCC AT AC A GAACAG AGGG GGAGGT GG GGG GT GT TA GG GC GTAG TT T G

File: 22CCRAA000_A11_premix.ab1 Sample: 22CCRAA000_A11_premix

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

Signal G:1900 A:1589 C:3303 T:3042

Lane: 95

Base spacing: -16.163063

1655 bases in 20860 scans

Page 2 of 2

620 630 640 650 660 670 680 690 700 710 720 730 GCTGA GGGAGAAA AG AAG CT GCC AGT T GAC GCGA GACA CGGCC AGC AGC AGC AGA A GACG CAGC AGG GGC GCTTCGTCTGGAGC T GAGGAA A AAGGG G G GGG C GAAG AT



