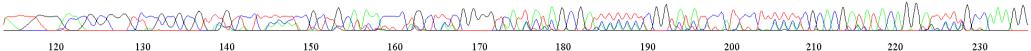


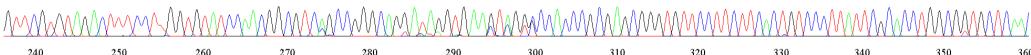
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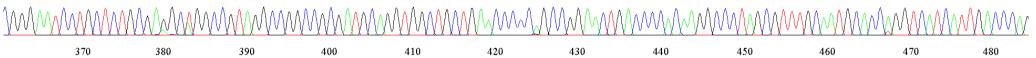
Sample: 22CCRAA000_G11_premix Lane: 83 Base spacing: 15.123373 1690 bases in 21181 scans Page 1 of 2

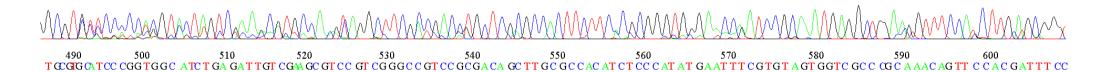
10 20 30 40 50 60 70 80 90 100 110 AATT GC TAA C C G GTCCCC GC GT T GF A CT T A A CG C TGCA A ATT C T G C T C G G G G T A T ATA G C G A TTTTTTC G GF AT A T C C T C T TTTTC C C GAT AT A CA G G A GTC T C C C C AAA G G



120 130 140 150 160 170 180 190 200 210 220 230 GTT CGT GT AG ACTT T CCTT GGTGT AT CCAA CGG CGT CAGC CGGGCAGG AT AG GT G AAGT AG GC CACC CG CG AGC GGGTGTT CCTT CTT CACT GT CCCTT AT T C GCACC T GG CGGTGCT CAA







File: 22CCRAA000_G11_premix.ab1 Run Enda
Sample: 22CCRAA000_G11_premix Lane: 83

610 620 630 640
T AGT C GATCA AGAC CTGG AA AAC GG GACGATT AT

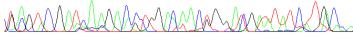
Run Ended: 2023/4/4 0:17:12

Signal G:2426 A:2747 C:4046 T:3327

Base spacing: 15.123373 1690 bases in 21181 scans

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Page 2 of 2



730 740 750 760 770 780 790 800 810 820 830 840 GTGTAC GG GCT ATA GAGAGA T C AC GC GATGCTATAA AC A AGT GG AAAAT GATGACC AT C GC GT C CAA GA GA T CGT GC ATCAA C G AA AACAT GT AGA CT GT GA AG AG T CAT GC GACTA

850 860 870 880 890 900 910 920 930 940 CGAC GT GGAT GTT TT GGG A A C AA AA C A AA A A C T G GAA G GAT AGT CCAT C G AAG G T T C G A G T T T TAT C C CGGGT C GG G G G A A T GAA C G CTC C T GG T G C

950 960 970 980 990 1000 1010 1020 1030 1040 105

G AG AT GGAAAACA GACCTTTTGTTT GGG GGAGGG CA ATT A CCCG GG TTTTTTAA T GAAT T GGGGCC C CGAA A GAAG A TG GGG CACA AC A A GGCC A A T GAA TTTTTTTT GG