

File: 22CCRAA000_H01_premix.ab1

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

Signal G:1695 A:2503 C:3453 T:2703

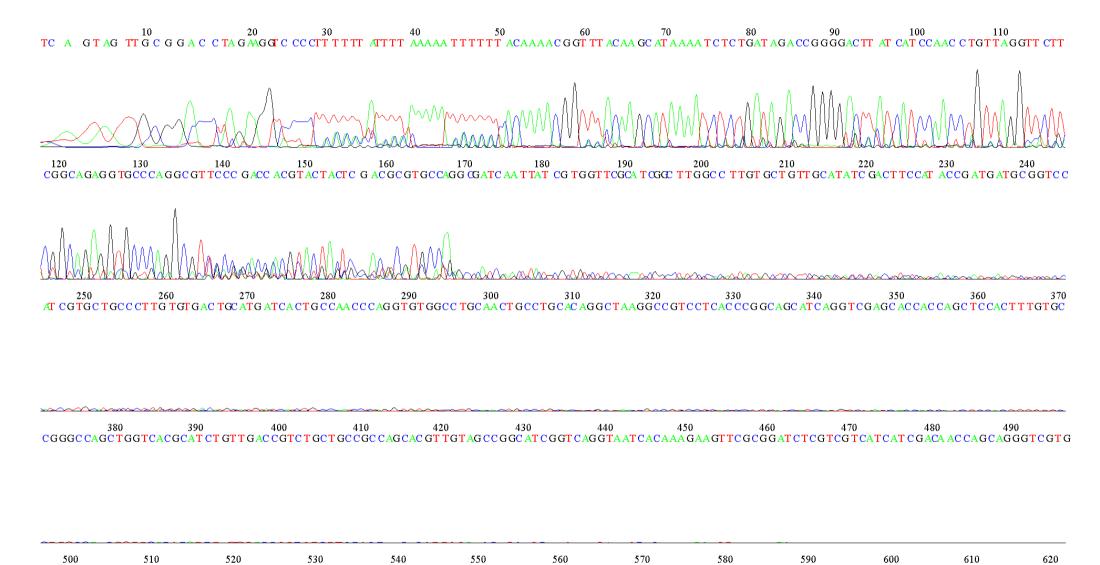
Sample: 22CCRAA000_H01_premix

Lane: 1

Base spacing: 15.520203

1085 bases in 16301 scans

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CTCATGCAA AGAA ACAC CTGTT CGGTGAGGGCCGGTGGCGCTCGTT ACGTT CAATT ACAT CCC CGT A CAT GT CGGGTCT GGAG ACCGG AA GGCT GAT C C CT AG AAT C CTAA CAAAAAAT CATC T G

Sample: 22CCRAA000_H01_premix Base spacing: 15.520203 Page 2 of 2 Lane: 1 1085 bases in 16301 scans 630 640 650 660 670 680 690 700 710 720 730 740 750 CA A AT AC GAA GCC GTC CCA A AT GAATC CCCCAAAGCA TC A CGC C CGTGT TC ATC C C C T TCAGT C A AAT GGACAAT T GC CT CAA GC C GCTGGACCTT CT GC CT T GC GT CGC C AT CA GCG GTGG TG C AG 750 760 770 780 790 800 810 820 830 840 850 860 870
TCC GAGCACG ACAGC A A CCCC TT G AACATC GAC G CCA CG CA GAT C GAA GACA G C GC A CTGAATG CCT GCCG A C GA T GCCA G CC A GGT GT GC TAT CC GT CG A G GA G CACAC GCA GCT CCAC T GGGC T 920 930 940 950 960 910 GGCAC TGAC GC G CG CA GAC C CGGC A T C GGTTACC TC CAT C AC C GC CT GCA GA GTG ATGGAA C GGC TC CT GCCCTACC A CACA CG C CG G CTAT GA A CCCCCCCCAGC G C TAC A CGAC C ATCAAT 1010 1020 1030 1040 1050 1060 1080 C ACAG AA T CGCTG CCCCCCAACAGAAAACA T GC G T TGT TC ATA T AC G A TT AGT GA T GC GA A TCGA A A AAC AC AC CGGAATAGT T C

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