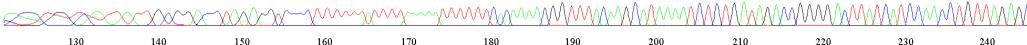


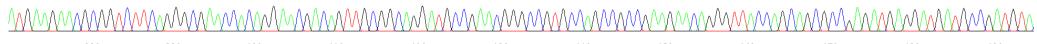
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10 20 30 40 50 60 70 80 90 100 110 120 G AAA TIT TOTA TA TIT GO G G ACCOTA G AGGI CCCC TITITTATTITAAAAATTITTTCACAA AAC G G TITIACAAG CATAAAAT CT C T G AT AG AC C G G G G ACTTATCAT C CAA C C T G T T AG AG





250 260 270 280 290 300 310 320 330 340 350 360 37
AT GAAGAACGCGGT CTT CAGGGCGAACCACAGG AACGACAGT T GCG GCAGG AT GCCGAACGGGCCGT GCCAGCCGCCGAAGAACA GGGT T ACCAGCAGCGCCGAGAT GAGGAT GAT GCCGAT GTA

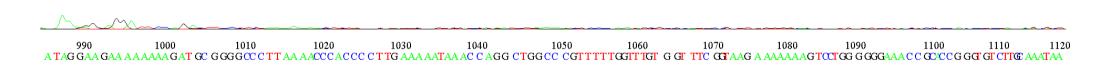


380 390 400 410 420 430 440 450 460 470 480 490 CTCACCGACGAAGAACATGCCCCACTT CATGCCGGCATACT CGATGT GGT AGCCGT CGGC CAGTT CCTGTT CTGCTT CCGGCT GGT CGACGGGT GACGGT GAT CACGG CG ACGC CAG CGAT GA



500 510 520 530 540 550 560 570 580 590 600 610
AGAAGGTGCAGAA ACCGAAGAACTG CGGAATGAT GAACCACAGGTTCTGCGCCTGGTACT CAACGAT GT CGCGCATGTTG AACGAACCCACCTG CACCACCACGCCCAT CAGCGCCCAG

File: 22CCRAA000\_H07\_premix.ab1 Signal G:1159 A:1479 C:2122 T:1287 Run Ended: 2023/4/4 0:17:12 Sample: 22CCRAA000 H07 premix Base spacing: 15.084803 Page 2 of 2 Lane: 49 1550 bases in 17735 scans GAA CACTT CGTAC GACACGGT CTGTGCCGAAG CACGCAAGCTGCCCAGCAGGC CGTACTT GTT GTT CGACGACCAGC CGG CGAA CAG CACCG CGT AGAC CGACAGG CC GGCCATGG C GAAGAA 750 760 770 780 790 800 810 820 830 840 850 860 G AA CAGCA GG CCCCA GGTG CCCA GGTG CG AT C G GGAT G AC CACG AA G C G AT TA GCA GG GGCT C A GGGC A CT ACC GGTGCCA GG AT G AA GAT C ACG CGTTC G A  $10^{10}$ A A A A GGGT AGGA TT CC A GAC TT CCTT GAAGAA A ACTTT TGA GA TTTT GAGGG GAAATT GAAA A GGCCC AA A GGGGCCAGT CCGGT TGGGGAACT A TGC ATCT A CGC GCA AC CCCCA AA GCGAG



ACCTCA C A ACTATGCCCTG GAAT GA GA CTCTAGGE A AT ATGA GAGAATA C AAAACGCTTGGGGGTTT TTAGGAAAGT G C GCC CT ACGCAACCAA A AAAACTGAGA GAAA GAGCGTC ATGTC C GCTGCGCGTCA T GCGGTT CCAT ACA