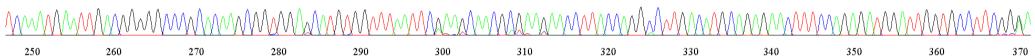


File: 22CCRAA000_G01_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:1182 A:1526 C:2058 T:1569

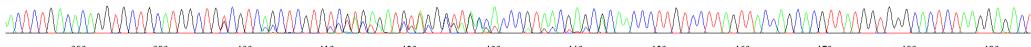
Sample: 22CCRAA000_G01_premix Lane: 3 Base spacing: 15.3786745 1680 bases in 18214 scans Page 1 of 2

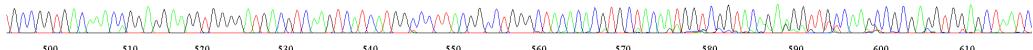


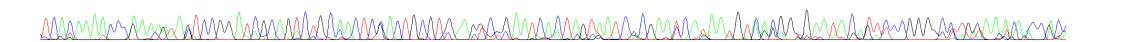
130 140 150 160 170 180 190 200 210 220 230 240
T CAAAT GAATT GACGGGGGCCCGCACAAG CGGTGG AGCATGTG GTTTAATT CGAAGCAACGCGAAGAAC CTTACCAGGC CTTG ACAT GCAGAGAACTTT CCAGAGATGGATTGGTGC CTT CGGGA



ACTCTGACACAGGTGCTGCATGGCTCTCGTGTCGTGAGATGTTGGGTTAACACCCCGTAACGAGCGCAACCCTTGTCCTTAGTTACCAGCACGCTTATGGTGGGCACTCTAAGGAGAGACT







Run Ended: 2023/4/4 0:17:12 File: 22CCRAA000_G01_premix.ab1 Sample: 22CCRAA000_G01_premix Lane: 3

Base spacing: 15.3786745

Signal G:1182 A:1526 C:2058 T:1569 1680 bases in 18214 scans

Page 2 of 2

C CGAT CAC AC CATGGG AAG AG GGTT GAA T CA GAA ATAA AA T AGT C GAG A CAAT A CGGAAG AA GGCTA AA CA C GGA GC C GTAT C AT G A T A GCTT T C GAA GAT AGTGCGA A GTAA AAA A GAG AACA A C





AT CT GAGTT AA AAA GA AA ATAAC GAA A GG AAC GG GGG ATGCGAAAA T GCAC CGC A TTA AG A ACTGCTG GA T AC ACGGACCAGAA TACAAGA T AGTG AAG A AG TCAGCTACAACCCG TCACT ACGTCAATCACCCG

