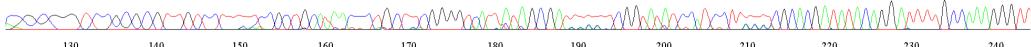
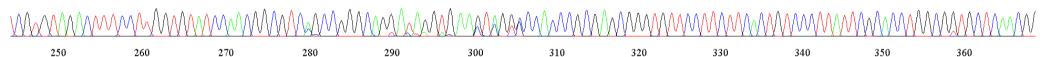


File: 22CCRAA000_F10_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:3088 A:3539 C:6145 T:4776

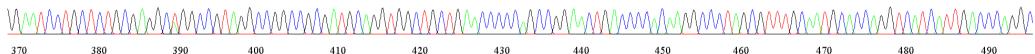
Sample: 22CCRAA000_F10_premix Lane: 70 Base spacing: 14.927772 1710 bases in 20560 scans Page 1 of 2

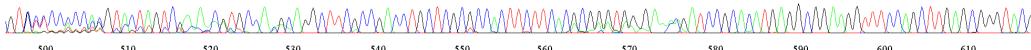


130 140 150 160 170 180 190 200 210 220 230 240 CGT GT AG ACTTT CCT TG GTGT AT CCAACGG CGT CAGCCGGGCAGGAT AG GT G AAGTAGGCC CACCCGCG AGCGGGTGT TCCT TCTT CACTGT CCCT TAT TCGCACCT GGCGGT GCT CAACG



GG AATCCTGCTCTGCGAGGCTGGCCGTAGGCCGGCCGCGATGCAGGTGGCTGCTGAACCCCCAGCCGGAACTGACCCCACAAGGCCCTAGCGTTTGCAATGCACCAGGTCATCATTGACCCAGGC





500 510 520 530 540 550 560 570 580 590 600 610 GGCT CCCGGTG GA GCT GA AAT A GT CGA ACAT C CGT CGG GC GT CGG GC GA CAGCTT CT CCCAA AT GAATT A CGT GT AAT GGT C GCCA GCA AACA GCA CGACGATTT C CT C GT A

File: 22CCRAA000_F10_premix.ab1

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

Signal G:3088 A:3539 C:6145 T:4776

Sample: 22CCRAA000_F10_premix

Lane: 70

Base spacing: 14.927772

1710 bases in 20560 scans

Page 2 of 2

CGA C AA GACCTOG C AACGGG ACGTTTT CTTGCCA COGTCCA G GAA GC COG A AGC GGTGCA TC AGC GACC CC CGATTC CAG GGGCCC A CACCOG TAC GACC GTAT C C GGT GG A CC CCAG TAT C CCAG TAT C CCAG TAT C C GGT GG A CC CCAG TAT C C



GAA G G GCG ACGGGC CT TT CCT CGGG CCATAAT AGGA GT AAAG GGT C C T GGATGCGA GA GGCT CGGGG T GT GG CAA AA GGT C CGG AAT ATGCGT AG ACG G GT CC ATC C A TCCGC C A GAGGATGGTTGGG

GTT GGT T GGAAAC AAAAA AAA GGAACC CAACC G G A GTT C C T GGT TC ATC G GATT T GT C AC C G CGA GGG GGAAG GAA TA AT CTTC GTGT TT GT A G AGGGTA AA AAGTC C A CCT C GGG GT GGG

GGG GCA ATT CCCAGCCGA GCTT TATTAGG TGA G CC CA AA AG AA TG GCAA A AAC C CCAT C AGT TAAGGT T GG GAGAA AGC C T C GGG GG GATAAA GCAC C GGT GC GGT GG GG CAT

A TGT T A AA ATA AACAG G A AAT GG GAAA AGA GGC TTT GG GGC TC T A TGTG G GGCGT C TCAAAT A CGG G TT GG GGG GAGAT TTTACACA A CAC CGC CCATG TAGT AAT TTCT T T A TGC CGA GC T TTA TATCG AG C