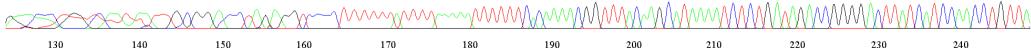
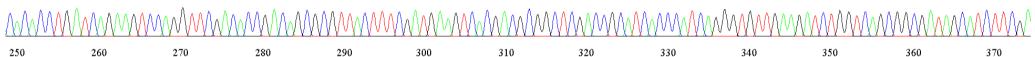
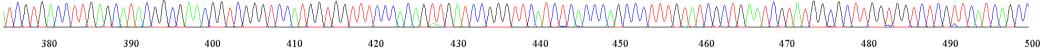


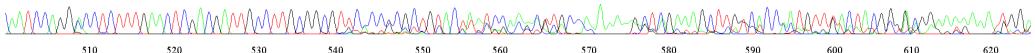
File: 22CCRAA000\_E08\_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:728 A:1065 C:1409 T:1081







AATC CAGGACCT CTTT GAACT CGAC GATT GCCTT GCGGCGTT CAC CGCGC TGT ACT AAT CCACG CTT GC CAAAA AA G CT CGAAA GCTT GT CAC CAAT GTT GACT A CG CGAAAAAA AAAATT GGA



510 520 530 540 550 560 570 580 590 600 610 620 C GAAAA AAAATCT C G GCTG AC A GT GAT GT GT AT G GC CACA AT AG A AAGAT AC ATT CT C C TT CT GAGT GAA AAA TGA G G G G G G G C T G T T C G CAGCA C T GT C GT GT T AAGAC G C CA A G

File: 22CCRAA000\_E08\_premix.ab1

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

Signal G:728 A:1065 C:1409 T:1081

Sample: 22CCRAA000\_E08\_premix

*Lane: 56* 

Base spacing: 15.001787

1804 bases in 22873 scans

Page 2 of 2

650 660 680 700 720 740 C C GCA CTTCGC A G A GGG AAAA A C AC ACACCA GG CA GTG GGAGCCGCAT GA GCG A GAA TAAT C T GCGT GC ATCGAT CGA GA CATGCT ACAT GG A CCGA CATA GA GA C CCG A GG A A GAA CTT GAT G



