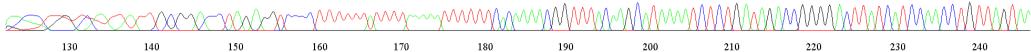
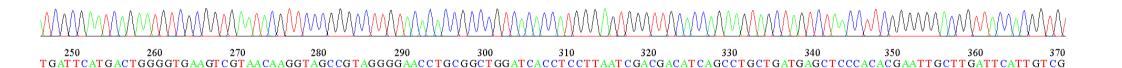


File: 22CCRAA000_G06_premix.ab1 Signal G:1252 A:1664 C:2233 T:1850 Run Ended: 2023/4/4 0:17:12

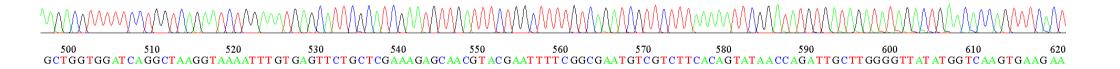
Sample: 22CCRAA000_G06_premix Base spacing: 15.171872 Lane: 36 1480 bases in 17635 scans Page 1 of 2

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380 410 420 430 440 450 460



File: 22CCRAA000_G06_premix.ab1 Signal G:1252 A:1664 C:2233 T:1850 Run Ended: 2023/4/4 0:17:12 Sample: 22CCRAA000_G06_premix *Base spacing: 15.171872* Page 2 of 2 Lane: 36 1480 bases in 17635 scans 630 640 650 660 670 680 690 700 710 720 730 740 GCGCAT ACCGTGG AT GCCTTGGC AT CCGTGG AT GCCTTGG AT GCCTTGG AT GCGTGG AT CCGTGG AT GCCTTGG AT GCGTGG AT GCCTGC AT ACCTTT GG GGAGT CA GCACA CAGA CTGT GAT CCAGAGATCT CTG AAT GG GGGAAC CACGC CAGC 870 880 890 900 910 920 930 940 950 960 970 980 990 AA GAAA CTCT ACCC TA CCAA AA AAA TAACTAGGTGGTACCTTG T GGGATTACCAT GGCCA GGTTTT GAA GA CC CCC GG AA AAA GA TTT CTG T G GA C GCC GACAA GAG G G A CC AT GA A GACA A 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110



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