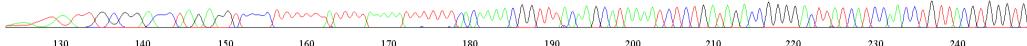


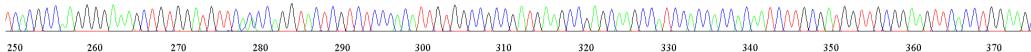
File: 22CCRAA000_H06_premix.ab1 Run Ended: 2023/4/4 0:17:12 Signal G:1842 A:2572 C:4018 T:2983

Sample: 22CCRAA000_H06_premix Lane: 34 Base spacing: 15.1399 1418 bases in 17529 scans Page 1 of 2

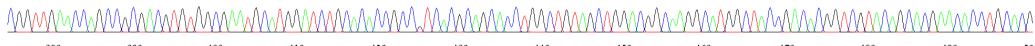
10 20 30 40 50 60 70 80 90 100 110 120 A A T AC T G C G G ACCCTA G AG GTC CCCTTTTTTTAAAAAATTTTTTTCCACAA AAC G GTTTTACAA G CATAAAAT CT CT G AT A GACC G G G G ACCTTTAT CAT C CCAAC C T G TT ACGT G G G T G

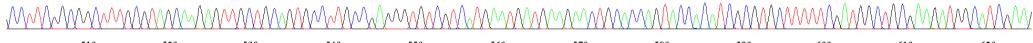


 $\frac{130}{120} \quad \frac{140}{120} \quad \frac{150}{120} \quad \frac{160}{120} \quad \frac{170}{120} \quad \frac{180}{120} \quad \frac{190}{200} \quad \frac{200}{210} \quad \frac{220}{220} \quad \frac{230}{230} \quad \frac{240}{220} \quad \frac{120}{120} \quad$



CGGTTGAACCAGCCGCGACGTGTGGCGAATGCATGGATCTGCACACCGCCGTCACGCAGCACCTGGCTGTAGCTGGACGCAAGGCATGGCTACCGACACGGTCGTACAGCACGAACACCTGCACG





510 520 530 540 550 560 570 580 590 600 610 620 GGTGGCTTTGCCATGAT CAGCAGTTATTTGCCAGGCAT GGCATTCGACCCAGCTT GGGCATGG CGCG CAAGGCTGT GTAC CTTTCC GACCT CT CGC AC GGT AA GGGCTT CTT CAA

