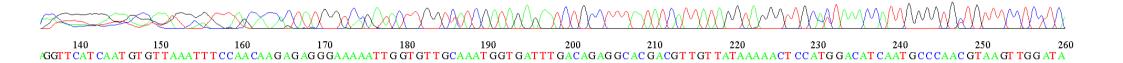
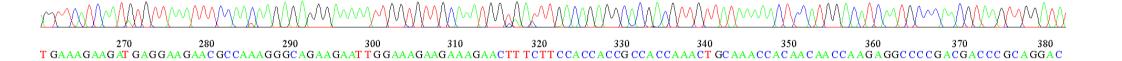


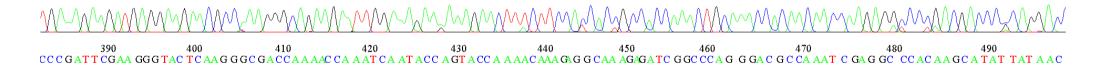
File: CYT2_CYT_R.ab1 Run Ended: 2025/5/10 3:56:54 Signal G:99 A:104 C:94 T:117

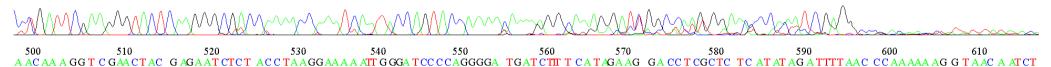
Sample: CYT2_Plate1_H11_CYT_R Lane: 81 Base spacing: 15.27879 2279 bases in 28088 scans Page 1 of 2

10 20 30 40 50 60 70 80 90 100 110 120 130 A CCTTCGCCAAAT CCCT AAATCC G TIT AGC A CCA C TAAT G GT CACGATACTIT G GT CTAT C GAT A GAG AATCT GC CT T TAA TAT CAT TAT A GGT A GGGT T GCAC T AAACCA T T G G GGT GCT GT T T CCT C C T C









File: CYT2_CYT_R.ab1 Run Ended: 2025/5/10 3:56:54 Signal G:99 A:104 C:94 T:117 Sample: CYT2 Plate1 H11 CYT R Base spacing: 15.27879 2279 bases in 28088 scans Page 2 of 2 *Lane: 81* TTA TAGGA GG GGAG A G AGA GATTCTGATA CCCC C CACTA ATT T GG A CACC C C A AAT CCAAGTTCA CT A TGT C TAT TCAGT A A AC CTCC A TG G CTTC T AACAGCAAT GACAT C T AC A TGCA C G C GTAA 750 760 770 780 790 800 810 820 830 840 850 860 870 TGCGT GCT GCT AT CAT AT T CAT AT T CAT ACT G A A CGACG AT A GT ACT G AT ACT CCTACT GCTGT GAT ACG AT ACG CG CT G GTA CTT CT GAA ACAA CCG A AT AT CGATGTACA GTCCACTGACT CG C A CGACGGATG C GATCA CTTCATG ATTCC GGT ACT AA CTAG TGACGC AA A AAAACA A CATTC GAATGT C T GGAATT A

