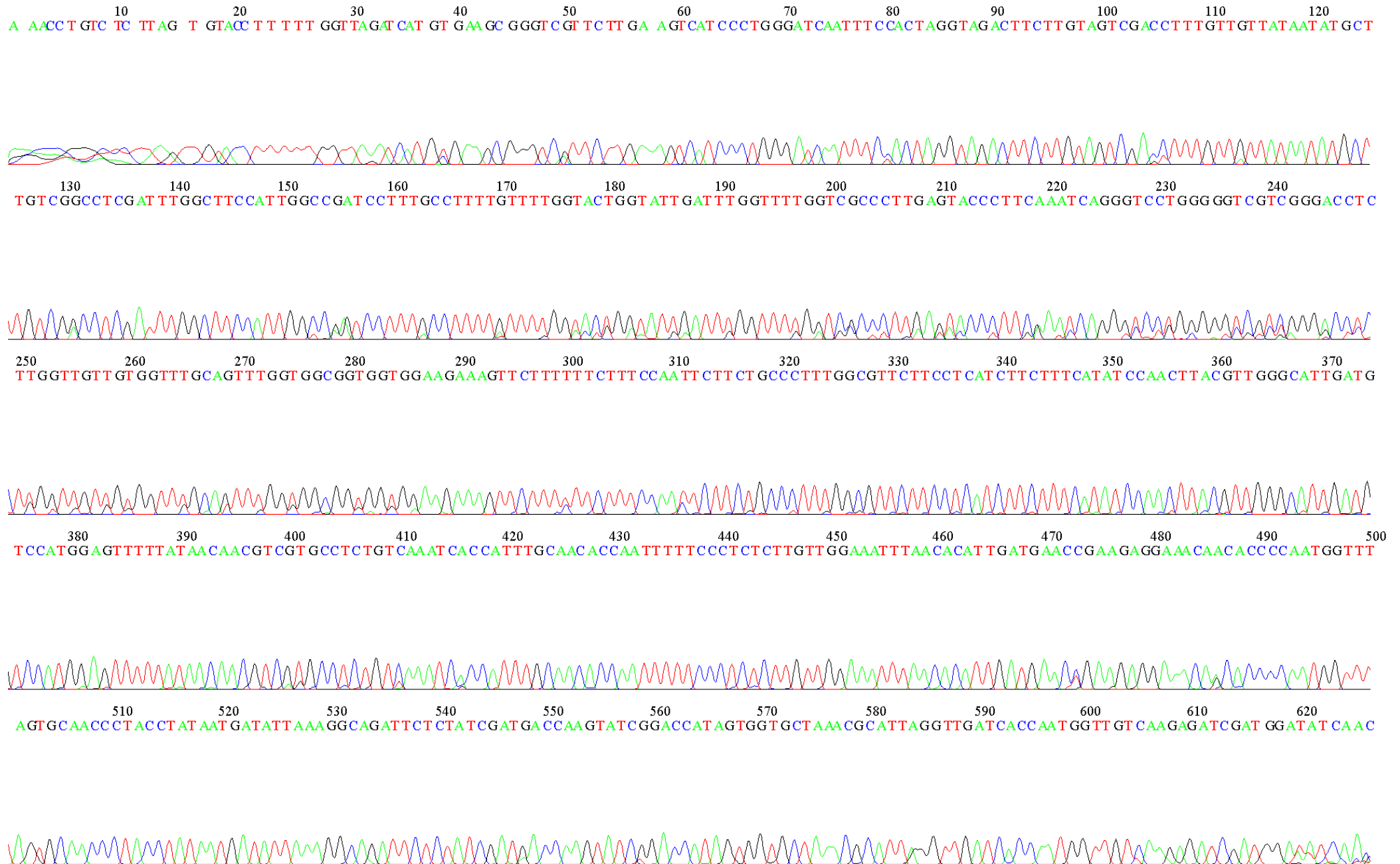


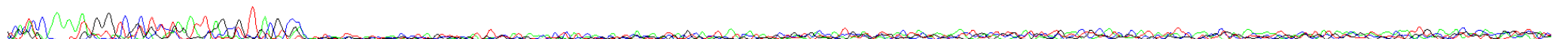
File: CYT2_CYT_F.ab1 Run Ended: 2025/5/10 1:59:13 Signal G:300 A:210 C:289 T:427
Sample: CYT2_Plate1_H11_CYT_F Lane: 81 Base spacing: 15.558864 1162 bases in 14002 scans Page 1 of 2



File: CYT2_CYT_F.ab1 Run Ended: 2025/5/10 1:59:13 Signal G:300 A:210 C:289 T:427

Sample: CYT2_Plate1_H11_CYT_F Lane: 81 Base spacing: 15.558864 1162 bases in 14002 scans Page 2 of 2

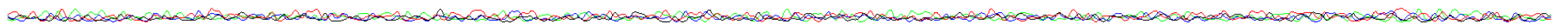
630 640 650 660 670 680 690 700 710 720 730 740
ATC AAA GGTCCTCTATTAGGTAGCCTTAT TTAGGCTTTGAACCCACGACAGAGATTAAGGCTA GACAACTATATGACTTCACATCTCA TTACAGATAAC AACATGAACCTA CAC



750 760 770 780 790 800 810 820 830 840 850 860
TAGTCAAATGTACTTCAGAA TCACTAA TCTTGA TCTACAGAGTATCTTGCTCTAGCTCAGTAA CCAACTAATAAA TTCAGTCAT CACA CCGGTGCTC ACTCTATCTTATATTTAAAC



870 880 890 900 910 920 930 940 950 960 970 980 990
GTAAAGAGAAATAATCTGAGTAA GTTTCGCATACAGTTTCGAACCTAAACAGTGGTATCTGAGTGACCCACAAACGAAAATC GTCTCAACAC TAA CACTGAATGTGCTGTTCGACTCCTCTGACTACACACGG



1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120
CTACTAAA GATCTTGAAATTCGTTCATGAACATTGGGTCACATCTTGA GCC TTCATATCCCCACTCCACTCTCTAGAGTT ATTAAGCGAAGTCTACTATCTCCTTGCTCCTTCGACACTTTACATCTA



1130 1140 1150 1160
TCTT CATGC AACGTGGCGTGT CAC ACTAAAA TTTCTG

