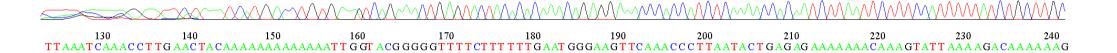
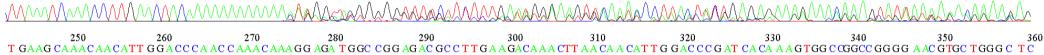
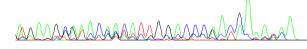


File: CYT6_CYT_R.ab1 Signal G:599 A:1116 C:1139 T:1303 Run Ended: 2025/5/10 3:56:54

Sample: CYT6_Plate1_C12_CYT_R *Base spacing: 15.43049* Page 1 of 2 *Lane:* 92 1732 bases in 21286 scans







400 410 420 C GGG A CTCCTCATC GG A ACCGC COG ACA GT CGAG CGG GTGG AGTC G G GC G ACCT A GAGA ACCT C AGGA GCAT CTGA CAG GTG CGA TTTTT C TGTT T C ACT G C CAGC GT G C G C CACG File: CYT6_CYT_R.ab1

Run Ended: 2025/5/10 3:56:54

Signal G:599 A:1116 C:1139 T:1303

Sample: CYT6 Plate1 C12 CYT R

Lane: 92

Base spacing: 15.43049

1732 bases in 21286 scans

Page 2 of 2

600 610 620 630 640 650 660 670 680 690 700 710 720 GGGTCCTC CT CCCTAGCATGAG TGCGC A CAACT CTTAAAT A A TGCTG GAG GGAGA AGGTAGA CC GTCTG C C GT GA G CAT TC T GTCAT TGT C A GATTGGTG G G TC TT CC GAC CAGCGACG C AC GGC C GAT CA T A

730 740 750 760 770 780 790 800 810 820 830 840 850 G AAAA CGGACC TGGGT CGTCGGG AGCTGT ACCT GCT GC AT ACT ACGTGTT ACCTGGT ATG C C C C AT ACAA GA CATGC G G GAGAAAAG C AC C A T A T

860 870 880 890 900 910 920 930 940 950 960 970
TA TGGAA A CCGTA CACAC CACTGC GTGGA T GCAC GG CGA GG C G A GŒAG CAT C CT AGATC TTCTAAC C A GŒC C AGCAA GG C T CCACCAT AT AA G C C CAG G GCAT CGTGACC AA CGCTA TA

980 990 1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 TTAG CATC AA ATGTAA GAT C G C CATC GTA TCGC AT GGA T C G C CATGTGA CAATCCT GTGACCG CCGC C G C G GAC CGTA CCAACTGGG A G GTAA CCG GA C G T A CAACGAG G GGA AT