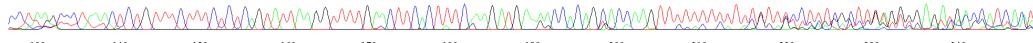
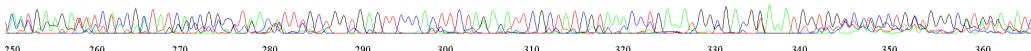


File: CYT6_CYT_F.ab1 Run Ended: 2025/5/10 1:59:13 Signal G:361 A:379 C:618 T:689

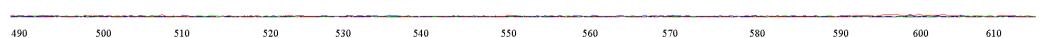
Sample: CYT6_Plate1_C12_CYT_F Lane: 92 Base spacing: -16.163063 3116 bases in 32767 scans Page 1 of 2







370 380 390 400 410 420 430 440 450 460 470 480
TGAAGCT A A A TGTTT GGGA T GCG A A GC C AT ACAT A CT C TTC GA CA AAT T CT G AGT GATAT A G CCT GA CACA G C T CT GG TG A CGC CAT CA CGAC GAG C AGC TT ATTIGCT A CT A CA CGAT AGTA



A CAAGAT TAGTAATG TGGT CGA C CAGAGT GT AT GAC GGATTGAA AT CGACTA GATTA CCA CACCAC A GCGC CCG CG ACTC CGCA T AAA G CG ACGCG CGT T GC GTT AA TAT ACGAAC CT GGTAA ACGT

File: CYT6_CYT_F.ab1

Run Ended: 2025/5/10 1:59:13

Signal G:361 A:379 C:618 T:689

Sample: CYT6 Plate1 C12 CYT F

Lane: 92

Base spacing: -16.163063

3116 bases in 32767 scans

Page 2 of 2

620 630 640 650 660 670 680 690 700 710 720 730 CGC T CA A GAACAATT CAAC CATCATGTT AT CAT AG A GTGAATGACTT AGCGAT CA AA CG A CCT GG GA CAT GAA GT GGT & AT GC A GGAA GGC CA CGATT GC AT CCT AC GC T GAT GAA CC

850 860 870 880 890 900 910 920 930 940 950
CTATGTATATTCCAATGTGCGCCGCACTAAGG GAAC CAATGA TATTCTGA CACCTC CAT CACTG G G T C CATTAT A T CA C TA A A A G A A TC TTG AAG C TA G AGCC G G C TC TA

960 970 980 990 1000 1010 1020 1030 1040 1050 1060
ATA AA AAAACTC G C C G T G G C A T T G G C T T A G C T T A A A T G G T G T T A T G T G G C T A G C G C G C G G G AAAC C A A G T C A C A T T

1070 1080 1090 1100 1110 1120 1130 1140 1150 1160 1170

FRA C. AAAAA A. T. C. AAACT G. G. C. T. G. C. T. C.

TCA G AA A T G A A AGT C C C GA A C G T C AG GTA CAT CT C T C GT GACT A GACT A AGACA AC ATGT ATG C AGC G A C T G G G A C C C AT G T G T TA C T C GTT AGT GA A GGA ACT