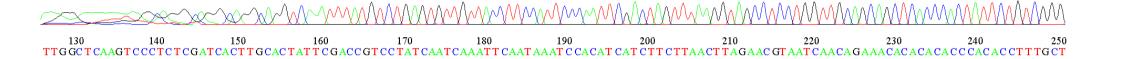
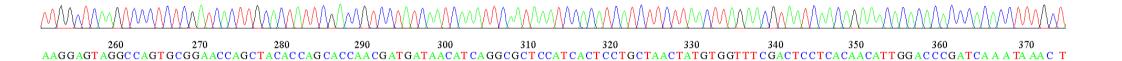
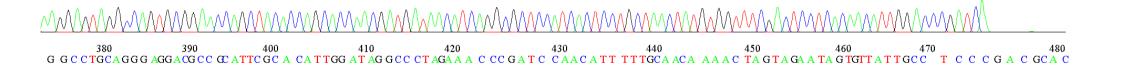


File: CYT17_CYT_R.ab1 Run Ended: 2025/5/10 3:56:54 Signal G:504 A:624 C:914 T:829

Sample: CYT17_Plate1_F12_CYT_R Lane: 86 Base spacing: 15.531295 1432 bases in 16841 scans Page 1 of 2







 File: CYT17_CYT_R.ab1

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

Signal G:504 A:624 C:914 T:829

Sample: CYT17 Plate1 F12 CYT R

Lane: 86

Base spacing: 15.531295

1432 bases in 16841 scans

Page 2 of 2

600 610 620 630 640 650 660 670 680 690 700 710 CTC A GGC G A A A G A AGTGA GT CCA GT TG TGC C AG A T AA T TGT A A T AC AAAGC ATG CTA C A A A C AAC AGGA A Œ GCA ATA GCAC ACTG CG G CA G GCCA C AGAGT GAG GC AGATGAAT GT AGA CA

720 730 740 750 760 770 780 790 800 810 820 830 840 ACC GTT A CAAGATGCTTA C GAT AT CTGAGA AT GCAA G TC GGCA CT C A G T GCT AAT ACA ACT AT AGCCC TT C TG G AT GG C AT G GT C AT C C G AG CT A GA CCA G AAAGCT CAAT GAATCA GTA G C T GAT AGC

850 860 870 880 890 900 910 920 930 940 950 960 970 C TA AA TGT TGGGA G CTGCCGTAGCTGAC GAA CAATC A ATG TTA A TAGGAGCCT GC G GTCC TT AGCAG TA CT TA T T TC AG A GCT GAT G TGCTA AG C G A AA GGAAG AT AACAT AGT T C C C TA GAA

980 990 1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100
TA GAA G G A A C GGIGAG A A A ACT ACG AG AC A ACGTCG GC CC AT GCCT T TAT CAAC CT GC T GITATA AT A C AA GT TGC TCACAA GT GI GITC TAC AAAC GAT AG T AAC AT T GCAT ATAT TAGT C C G T GIT ACT A

1110 1120 1130 1140 1150 1160 1170 1180 1190 1200 1210 1220 1230 1240 1250 1260 A T A C AAT A CGGT C GGTGT CACCCGTGCGCAGGTGCCAGCAT A C GCATGA CTCAATCAT GTAAC CT AGA AT AAT GGGT ATATTCGATGACGA AATAACATA TA CACTT G AATG ACTTT AGFACGT A TG CGGC GT CACGCA CTGTGTGT GC