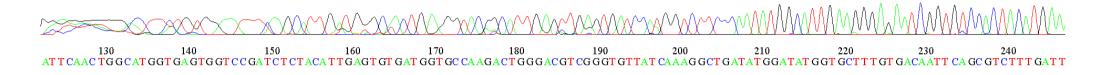
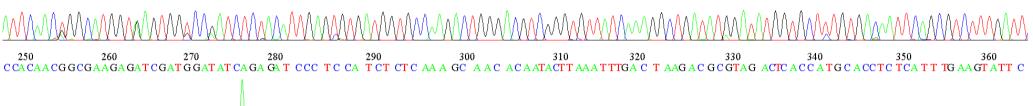


File: CYT5_CYT_F.ab1 Run Ended: 2025/5/10 1:59:13 Signal G:463 A:381 C:378 T:508

Sample: CYT5_Plate1_B12_CYT_F Lane: 94 Base spacing: 15.645171 3320 bases in 30181 scans Page 1 of 2

10 20 30 40 50 60 70 80 90 100 110 120 TCT GCCT CAT AC TT CTG TG G A TTT CA GCATT GAT GT GT GCTACT GAG GGT AC GGC ATT G AGTT GAGT GAT CCAT G GT CTT CCCA AT ATT G GCATT G AA GGG AG AAT C GGT GT CCAC T A CAT G AA







490 500 510 520 530 540 550 560 570 580 590 600 610 620
TIGT CA ACITT GT GAAC CTCA C ACAC G TTG CACA GOATT A AIT AC ACTT AAACT CAT CAT A AAAAT CACTC AG A AATAC ACOCAA TAT GG T CA T C C GIT CAT C C AATA G GT AAGA G G T GAT A AT TAT

File: CYT5_CYT_F.ab1 Run Ended: 2025/5/10 1:59:13 Signal G:463 A:381 C:378 T:508

Sample: CYT5_Plate1_B12_CYT_F Lane: 94 Base spacing: 15.645171 3320 bases in 30181 scans Page 2 of 2

630 640 650 660 670 680 690 700 710 720 730 740 750
TCGC A ATTTATGT ACAT A CT C TCTTATA A AATCAT GACC ACT GAAACA CTT A T CATA A CCATACATA CTT CA ACCG CCCACGTAACT AT CCT CT ACAGAGAGT C AA T GCGTCGAA CACT AT CATT T ACAAT A C T T G

760 770 780 790 800 810 820 830 840 850 860 870 880 890 900 ATT TCCGCGACGCTGTGAGC GGTTCT AT C GCT AT GT ACTA ACTT ATA C ACATCGA A AT TGTTTAT C TTATT A A A T G ATCTCGCC ATTCA ATTTT AT CTA ATA GATTGTTG T ATA C TGTTGC TGC TGCGACC

910 920 930 940 950 960 970 980 990 1000 1010 1020 1030 1040 T TGTAÆTA GCAT ACGAT GTAT TAGAT TAGAT TATGATCTT AAGC CA TACT CGT AGAAC AG ATÆGTGA CT ATC TTGÆT GCAT GCGAGT GTGGT GCGAT GTCAACT G

ATGT CGCCTTAC A CC CTGT CT GGCAC AGT CACA T GATT AT CGT CGCACCGA CGTTGT GT TGC ACTGC A CAC TG AT GT ACATTGAC CGCCGAT AT CT CCGTGGGATGTT C GT GT CC AT CAGA AAGTACT CC GA CT ATCCCCGTGC AT A