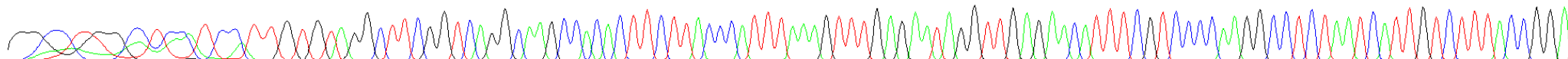
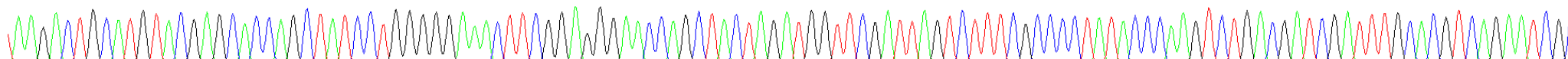


File: 21-MI-1_D3-D5Rv_premixed.ab1 Run Ended: 2022/6/9 16:39:58 Signal G:1325 A:1667 C:2613 T:2185
Sample: 21-MI-1_D3-D5Rv_premixed Lane: 32 Base spacing: 15.69409 1062 bases in 21012 scans Page 1 of 2

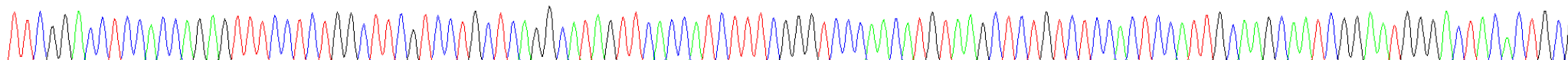
10 20 30 40 50 60 70 80 90 100 110 120
T GAA ATGAA C T AA T CCAT T GT GTAGGCTTCG GT CAGGCAAGCCACACTTCTTACCCATTTAAAGTTTGAGAAATAGGTTGAGAACATTTCTGCCCAAGGCCTCTAATCATTGTCTTTACCGGAT



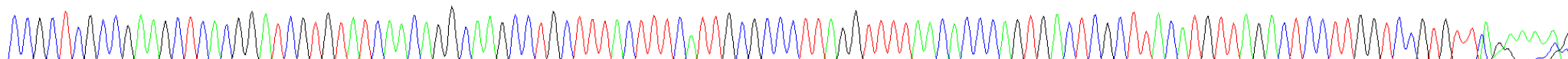
130 140 150 160 170 180 190 200 210 220 230 240
AAGACTGCATGTACGAGCACCAGCTATCCTGGGGGAACTTCGGAGGGAACCAAGCTACTAGATGGTTTCGATTAGTCTTTCGCCCTATACCCAAGTCTGACGATCGATTTGCACGTGAGAATCGC



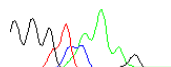
250 260 270 280 290 300 310 320 330 340 350 360 370
TTCGGACCTCCACCAGAGTTTCCTCTGGCTTCGTCCTGCTCAGGCATAGTTCCACCATCTTTCGGGTCCCAACATGTAAGCTCTGTCTCCACTCCATTGCAAGCAATCGGAATGGGACTACACTGCG



380 390 400 410 420 430 440 450 460 470 480 490 500
CCGCTCGCCGAAGCTCACGGATCGTGTATCAACAGGCAAGCCTGCTTTACTTTTATTGCGCCCTTAGGTTTTAACACCCAGTGACTCGCTTACATGTTAGACTCCTTGGTCCGTGTAGAAAAAG



510 520 530 540 550 560 570 580 590 600 610
GGGTAAAGAGGATTCTAAAAAGGTAAGCGAGTCA GTGGGTGTTAAAACTTAAAGGCGCA TGAATGTAAAGCATGCTTGCC GTTGATACACGATC GGT CGCTTCGACTATGATGCAATG



620 630 640 650 660 670 680 690 700 710 720 730
TAA T C CAT T CT AT AG CTA GC AAT GG AT T T G G AGAC GCAG C T T ACATG TGG G G A CCGA A G A T T G T T AAATAT G CTGTAGC CGCAGTAAT CGAAGGT AAACTCT G C T GGGAGG TCCTAA

740 750 760 770 780 790 800 810 820 830 840 850 860 870
AC A TA CT GACCTGC AAA TCG AT T GGT C GTACGTGG GCTACTGG GGTGCGT AAA GAAC TAAATAC CCACAT CATGTTCA AC TC GATTTCCATC TAAA TTT T C C C C G G A A G G G C G G T G CT GG T AAAG

880 890 900 910 920 930 940 950 960 970 980 990 1000 1010
GGCCA CTTCTT TAC T C GGTTCGAA CAAAAATGG AATATAAAA GGGCCT T GGGGGGA AA CAAAGTAGCTTTC TA CCTTA TTC ATCAACGCT T T T AAAATTG G GGT T AAAA AAAATTGGGG C C C T C T G C C C G G A G

1020 1030 1040 1050 1060
CCGAAG C C AACC CACATGT GGAATATTTT TAGT GTG CCC GGTGGG