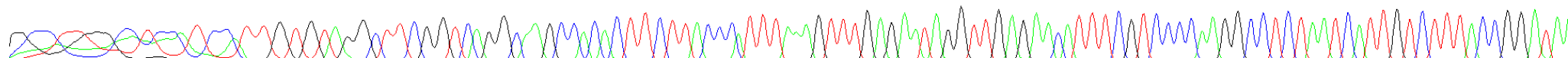
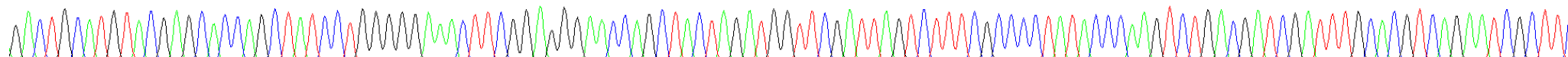


File: 36-MI-2_D3-D5Rv_premixed.ab1 Run Ended: 2022/6/9 16:39:58 Signal G:958 A:1200 C:1845 T:1578
Sample: 36-MI-2_D3-D5Rv_premixed Lane: 24 Base spacing: 15.524863 995 bases in 16614 scans Page 1 of 2

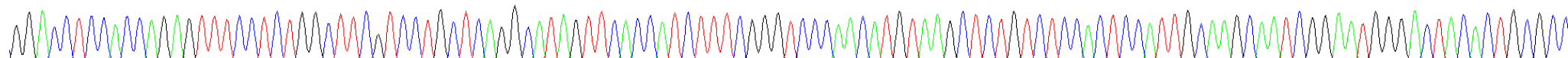
10 20 30 40 50 60 70 80 90 100 110 120
GGCAATAGG C TCCA T CCA T T G T GTAG GCTT CG GT CAG GCAAGCCACAC TT CTTA ACCATT TAAAGTTT GAG AATAGGTT GAG AACATT T C GT CCCCCAAGGCCTCTAATCATT GT CTTTACCGGATAA



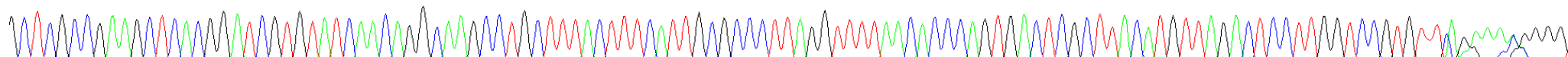
130 140 150 160 170 180 190 200 210 220 230 240 250
GACTGCATGTACGAGCACCAAGCTATCCTGGGGGAACTTCGGAGGGAACCAGCTACTAGATGGTTTCGATTAGTCTTTTCGCCCTATACCCAAGTCTGACGATCGATTTGCACGTCAGAAATCGCTTC



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



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



510 520 530 540 550 560 570 580 590 600 610 620
CAA GAAAGGATTCTAAAGGGTAA C GAGTCAGTGGGTGTTAAGACCTAA AGGCGCAGTGAA GTAAAGCAA GCTTGCCAGTTGATACAC C TCCGTGCGCTTC G T A GGG C CAGTGATAGTCC



630 640 650 660 670 680 690 700 710 720 730 740
GG T C G TAGGCT G C AAGGAATGG AGACAG A GCTCAC A G TTGG G AC C G AA G AT GG TG AAATATG CCGAACCCCGACGAATC GTACGAAA CTCTAGA TTG AAG T CC AAACGT TGCTCA ATGCA

750 760 770 780 790 800 810 820 830 840 850 860 870
AAT GA TG T C GAAATG G C TATT GGGGGG GAA AG ACCAATC GAA CACCTCA TAACT G GTCC TC AAAGTTT CCCGGAAC AC GGGGGG C TG A ATGACAATCTT TCCGTTA AAAAAACAT A AAAAAAGGGC

880 890 900 910 920 930 940 950 960 970 980 990
GT GG GGAA AAAATGTTTTCCTCCAT TACTA AGAGTT AAAT GGGTTTAAAAA TGTGG GCTGGG CC G CCAGG GC C CCCAT GG AATTT T GA TAGGCGGAT A AACCTTTT T AGGTG CAG AAA