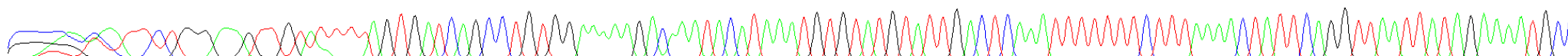


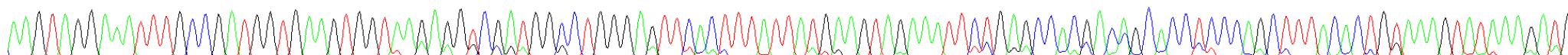
File: 1E70ZAC098.ab1 Run Ended: 2020/5/14 22:59:2 Signal G:2203 A:4238 C:3767 T:5075

Sample: 1E70ZAC098_premit Lane: 66 Base spacing: 14.626461 2131 bases in 24485 scans Page 1 of 2

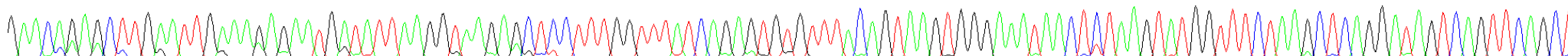
10 20 30 40 50 60 70 80 90 100 110 120
A CGTAG TA GTC TGG A GTT GTATTTTAGT GATCAGCC TGTGGAAAAAGACAAATACATAAATGTGTATGTATTTGACTCAAATTTTTTTCTTTAAAACTATTTCAGGTTAATTATAGAAAATGCC



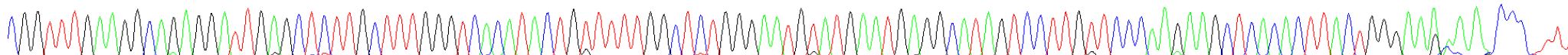
130 140 150 160 170 180 190 200 210 220 230 240
AAGTAGGAAATTTGCCGATGGTGAAGTGGTAAGAGGTGCGATGGCCTGGGAGTTCACCTTTATTATGAAGTAGAAATTCAGCCACGACAGCACCTCCCAGCTTTACACTGTAAAGTATAAAGATG



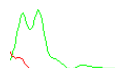
250 260 270 280 290 300 310 320 330 340 350 360 370
GAACAGAGCTTGAATTGAAAGAGAATGATATTAAGGTAAGAGCTCCTTTGGTTTATCAGAGTGTGTTTACAGTAAGTGGGAAATAACTCTAAGTATGGTTTCTAAGCTCAGGATAGTCAGTTCAGCT



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



510 520 530 540 550 560 570 580 590 600 610 620 630
AA A A AAAAAAAAAATTTCTTTAAAAAAAAATAGGTTAAATAAAGACCTTCCTTGGGGGGGATGGGGGGGGGGGTGAGGGGGGGGAAAAGCCCCCAAAACAAACGCCCCCCCTTTAAGCTCAAGGGATGAAAGGG



File: 1E70ZAC098.ab1 Run Ended: 2020/5/14 22:59:2 Signal G:2203 A:4238 C:3767 T:5075

Sample: 1E70ZAC098_premit Lane: 66 Base spacing: 14.626461 2131 bases in 24485 scans Page 2 of 2

640 650 660 670 680 690 700 710 720 730 740 750 760
GTGTGGGGAGGGCGAGCGGCTTCAAGACCGCAACGGGTCTGGTGGTTGTAGTCCCCCGGCCCTTTTGGGGTTGCGTGTTTTTTTGAATAGTTTTCAAGTTAATTTTTTCAATTTTTTAGTCGCGTGGGGCCAAC

770 780 790 800 810 820 830 840 850 860 870 880 890
AATTTTTATCAGGCCAAAAAATAAATTACTGGACATATAAACATTTTTTGAATACAAATAAGATTGATTAGGTTTATGGGTGTTTTTAGGTCAAGAAAGAGTTGGTAAAATGGAATTCTGTGTTATG

900 910 920 930 940 950 960 970 980 990 1000 1010 1020
GGTAGGGTGGAGGGTTTGTGGTGAACCAAGTGGGTTGGTTAGGAAGGATGAAAGAAGGGAACCTGGGGGGGGGATAGAAAGAGTAGAGTTTAAAGGGGTATGAGAAAGGATGTGGTAGGGGCGATAAGAG

1030 1040 1050 1060 1070 1080 1090 1100 1110 1120 1130 1140 1150
GCCCCGAGTGGGCGCCAGCACCAAGCGACCGACGCTGAAATAGGAAGGGTGAGCCAGTAGATGCCAGCCACCCCGGGTGGCGACAAAGACGGAAGGGGTGGACGGCAATCGGCTGCGGGCGAGG

1160 1170 1180 1190 1200 1210 1220 1230 1240 1250 1260
CGAGACATATCGAAACTGCTTAGGGCAATGGGTAAAGAACGTTCTGCGAAACGGCGTGGCGTATTGAGGAGAACGGCAAAATACTTCCAATCGCAAGGCTTGGGGCCCCCTCCCGCTAG