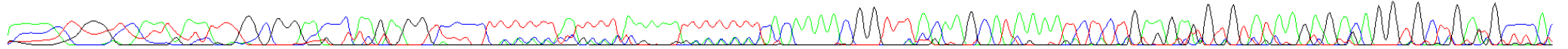
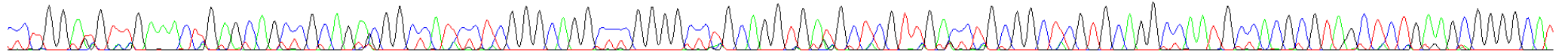


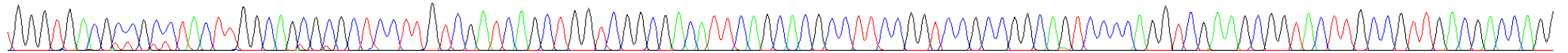
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T A A C T G T C A A G A A T T T G C G A A C C A A G A G G T C C T T T T T T A T T T T T A C A A A A C G G T T T A C A A G C A T A A A A T C T C T G A G G A G G T G T A T A A G A G A G G T G C T G A T C T T A C



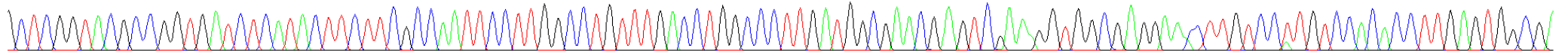
130 140 150 160 170 180 190 200 210 220 230 240
T T G G A T G T T G A A A C T G A G C A G C C G C A G A C G G C C A T T T T C G G G C A G G C T T G G G G C T T G C A G G T G A T G C T C G T T G A C T T G C G G C T C G T G C A G G C C A A T G C C A C G C G T A G C T C T G A A G C G G G G C A T



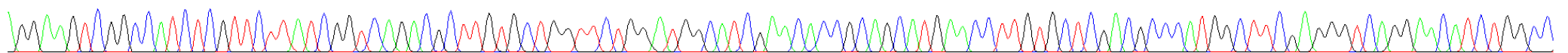
250 260 270 280 290 300 310 320 330 340 350 360
G G G T G A C G C C G C C T A C T T G G C A G C G C G C T C C T T G T C G A T C A G C T G G T C G G C G A C A T T C A G C A C G C C T T C C G G T C C G C G G C A G T C C C C A G G T C G A A G C G G T A C T T G C C G T T G A C G A C C A G



370 380 390 400 410 420 430 440 450 460 470 480
G C T C G G T A C G C C G G T G A T C T C A T A C T T C T T C G C C A A T T C C T T G G C C T G T T T G A C C T G G C C C T T G A T G G C G A A C G A G T C G A A G G T G G C G A G G A A C C T T G T C C T T G T C C A C A C C T T G A G T G G C G



490 500 510 520 530 540 550 560 570 580 590 600 610
A G G A A G T C G G C C A T C T C G T T C T T A T C G G T C A G A C G C T T G T G C T G G T T C T G G A T G G C A T C G A A C A C C G C A G C A T G A A C C T T G T G T C G A C G C C C A T G G C T T C G A G G G T C A G G A A C A T C T G G C C



File: 2103ZAB014_premix.ab1

Run Ended: 2022/1/28 0:49:8

Signal G:4773 A:6010 C:9740 T:6757

Sample: 2103ZAB014_premix

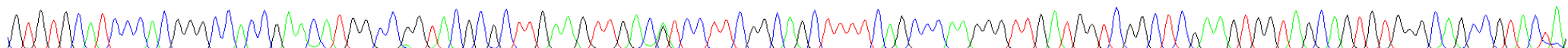
Lane: 36

Base spacing: 15.113358

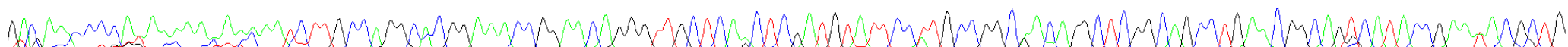
1474 bases in 18770 scans

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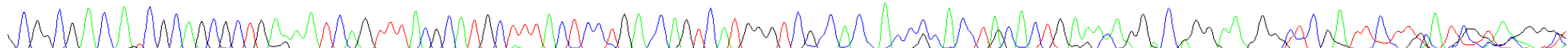
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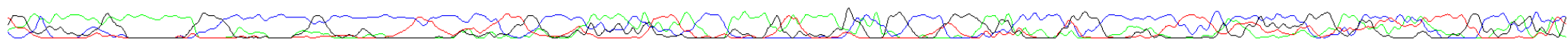
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860 870 880 890 900 910 920 930 940 950 960 970 980
GCGGCG A CTAGCGCAGCGCTGAGAA T CAGTTTACGCATGCTTTACTCCTGACCAGTCATGGGTGCGCA CCAC CCCACCTACACTGAAA AGGACGGGAA GGGCTC GATTC TTC ATCG TACGGC



990 1000 1010 1020 1030 1040 1050 1060 1070 1080 1090
G AAA A AGGAAAA AGG CCCCCCGCCG CCCTTCCCCCTTGCC C AAA CATTA C C AA AGAAAA GAAGGCC C GGAAACCCCGGGCC ACC CTC CGC CCCC G GAACCTCAATTTC C C A C A



1100 1110 1120 1130 1140 1150 1160 1170 1180 1190 1200
G GGG GCC TTGGAAGGGTTT G AAACCC GGGGC G G C C C C C T G C G G A G T G G G G G G G G C C C C C A A G G G C C C C A C A A A A A A A G A A A A A A G G A A A A A C C C C G A

