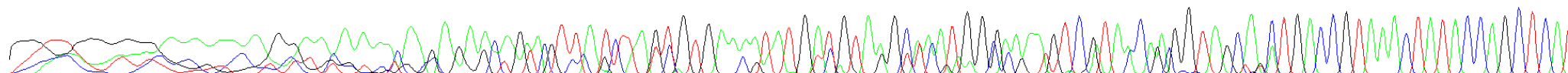
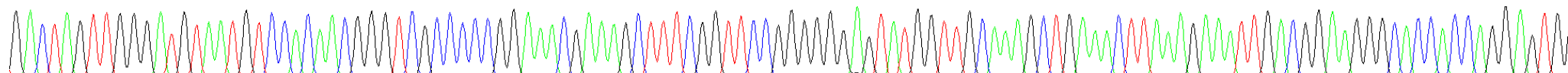


File: D3\_1813F\_premixed.ab1 Run Ended: 2022/6/9 16:39:58 Signal G:651 A:1115 C:956 T:940  
Sample: D3\_1813F\_premixed Lane: 56 Base spacing: 15.574707 1367 bases in 16804 scans Page 1 of 2

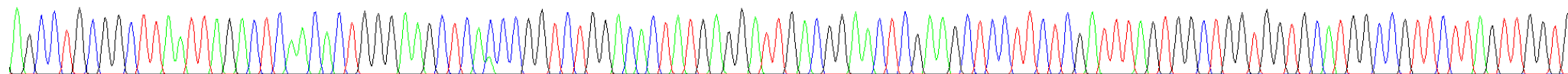
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GG CT CT G ATAT C A AG TACA TG GAA C G AAACACG AGA ACAG A GTT A TCTA GT GT G AAAAT AT GAT GT AA GCAAAT GGCAA ATGT C GTACCAAGGT AGCGCT GACCGTAAACTAT ACCAGCT CAT



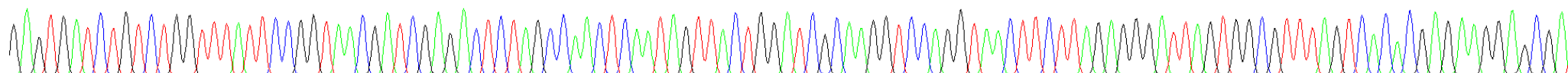
130 140 150 160 170 180 190 200 210 220 230 240 250  
G A C T A G T T G G G A T G T A A T G T C C A C A A C G G G T C G C C C C C G G A A A C G A A A G C T T T C G G T T C C G G G G G A G T A T G G T T G C A A A G C T G A A A C T T A A A G A A A T T G A C G G A A G G G C A C C A C C A G G A G T G G



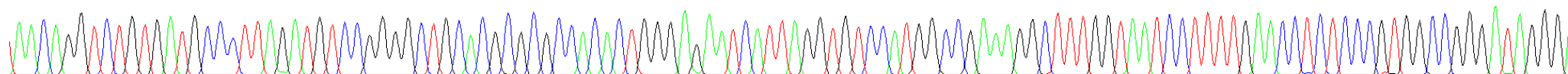
260 270 280 290 300 310 320 330 340 350 360 370  
A G C C T G C G G C T T A A T T A G A C T C A A C A C T G G G A A G C T C A C C C G G T C T G G A C A C T A T G A G G A T T G A C G G A A C T C G A A G C T C C T T C T C G A T T T A G T G G C T G G T G G T G C A T G G C C G T T C T T A G T T G G T G



380 390 400 410 420 430 440 450 460 470 480 490 500  
G A G T G A T C T G T C T G G T T T A T T C C G G T A A C G A T C G A G A C T C T A G C C A A C T C A A T A G T T A C T G G A T C G C A A G G T C C A G G T A A C T T C T T A G A G G G A T T A G T G G C G T T T A G T C A C A C G A G A A G G A G C G A T



510 520 530 540 550 560 570 580 590 600 610 620  
A A C A G G T C T G T G A T G C C C T T A G A T G T C C G G G G C T G C A C G C G C C C A C A C T G G G A G A A T C A T T A G G T G T C C A T G G C C G A A A G G C T T T G G T A A T C C T T T T G A A C C T C T C C C G T G G C C G G G A T A G G G A



File: D3\_1813F\_premixed.ab1

Run Ended: 2022/6/9 16:39:58

Signal G:651 A:1115 C:956 T:940

Sample: D3\_1813F\_premixed

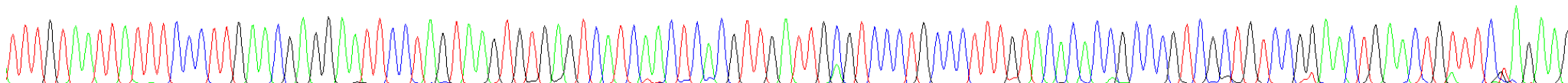
Lane: 56

Base spacing: 15.574707

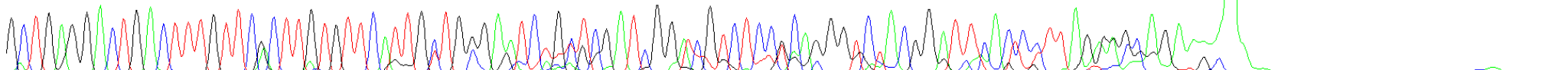
1367 bases in 16804 scans

Page 2 of 2

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760 770 780 790 800 810 820 830 840 850 860 870  
GCTGAGGACTGACTTTGTTGCTTGTGTTTCATTGCTGGGAATCTGCTCGATCGGACGTCTCCGCA GGGTCTACGGATT CACCCTTAGAAGCAGAAG AAA AAC TCC A ACCACG CC CACCC CC



880 890 900 910 920 930 940 950 960 970 980 990  
CTCGGG C GGGG AG C G T ATGAGC GAAAA GTG A T G AAGAA AAT AGGA CAAA GGTGAAGGGCCGAGTTGTG G GAC GGATC TAGG ACGTG ATAGAA TATG GAAAG GATT C TAGGCCCGT GGCAC



1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100  
CC TGATC G G G A T TACAGTG T T G T ATAGATGG T TGT GA TTAT GTTGT CAG CACGG A GGG TTAG AC ATTAATT TGCTTACGTA T T GAT AC A GTGT ATGCG GGTG CTACGG C GTTAG



1110 1120 1130 1140 1150 1160 1170 1180 1190 1200 1210 1220  
TCTCATGTCTCGGGCA GAGAG AA CG TATGGG GAGG TGGACGG GGGAGG A C G AT CAG TG TG AGT GTA TG ACACGAA ATAATAA G T GGGG T GGA CA G GATTTGTAGG T C TGT CGGATGT

