

File: 21-MI-3_D3-D5Fw_premixed.ab1

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

Signal G:1703 A:2341 C:2478 T:2562

Sample: 21-MI-3_D3-D5Fw_premixed

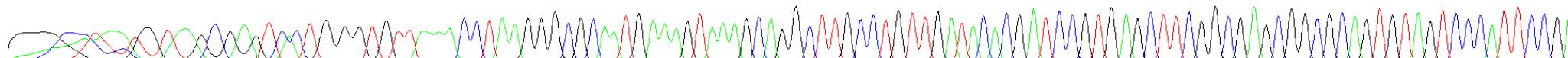
Lane: 11

Base spacing: 15.615305

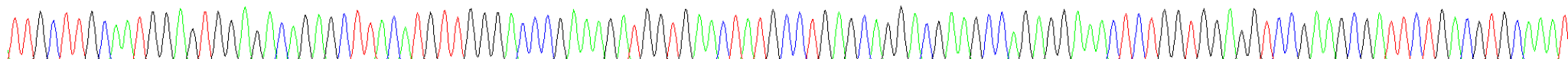
1802 bases in 21045 scans

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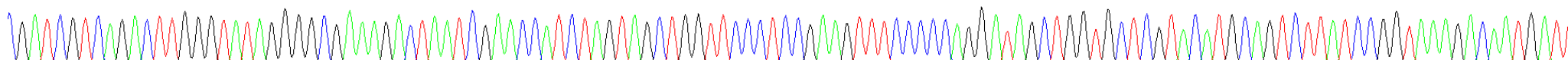
10 20 30 40 50 60 70 80 90 100 110 120
A GACCTAC TG TAGC GAGTCCT G GGT GTTAAACCTAAGGGCGCAATGAAAGTAAAGCAGGCTTGCCTGTTGATACACGATCCGTGAGCTTCGGCGAGCGGCGCAGTGTAGTCCCATTCCGA



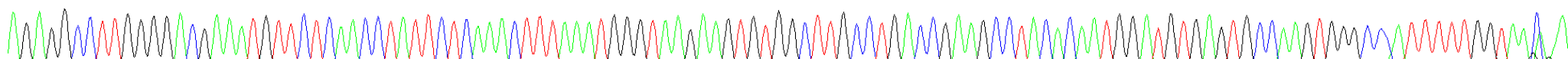
130 140 150 160 170 180 190 200 210 220 230 240
TTGCTTGCAATGGAGTGGAGACAGAGCTTACATGTTGGGACCCGAAAGATGGTGAACTATGCCTGAGCAGGACGAAGCCAGAGGAACTCTGGTGGAGGTCCGAAGCGATTCTGACGTGCAAAAT



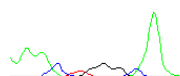
250 260 270 280 290 300 310 320 330 340 350 360 370
CGATCGTCAGACTTGGGTATAGGGGCGAAAGACTAATCGAACCCTCTAGTAGCTGGTTCCTCCGAAGTTTCCCCAGGATAGCTGGTGCTCGTACATGCAGTCTTATCCGGTAAAGACAATGATT



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



510 520 530 540 550 560 570 580 590 600 610
AAACTGACACTTGCTCGTCTCGTGTAGGCGTCTCGTCAAGG C AGCCACTTCTCACCATTTAA GTTGA GAATAGGTAGAGAACATTTCGTCCAGGGCTCTGATCAATGTCTTTACG



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620 630 640 650 660 670 680 690 700 710 720 730
GTAGGACAC GCAGGTACGAGCACAGCTATACAGGGG GAACTTC GATGGGTACAGCTCTAGATAGTTTC ATGGAGTC TTTTC CCGTGTGCCCGGGTCTGAACAAATGGATCTTGCAAGGTCG

740 750 760 770 780 790 800 810 820 830 840 850 860
AAATGCTTTC GAAC TCCCCAAA AATTTCCTTTCCTGGCTTTCCTC GGTAAAGGGGCAGAA TTTCAA TCTCTTTTGGGAGGCCCAAAGGTA AACCTTTT TTTTCTCCCTCCCTTTGCGCGGCGTGTCAA AAAAGG

870 880 890 900 910 920 930 940 950 960 970 980 990 1000
GAGGAAACAAACCGGCCCGCCCTCC TCCACCTATCAGACACATAAGAAAGGGGGGAAGAGACGAGCCGGAGCCC C GCTCTCTCTTTTGTATTTGGCTGCTGT TTTGTTTGTATTTCACCCACCCCGGCGGC

1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120 1130
GGCCGAGTGATATAG ATACCATCGGGGGG GTGGGGTTGGCTTAGGAAA AAAAGAGAAAAAAA AAAAAAA CCAACAA GGGTGGGTAGCGGATCGCTTATGGCTA GGAGAGGACATCA

1140 1150 1160 1170 1180 1190 1200 1210 1220 1230 1240 1250
GCACCATTCACGTGC GGCC TGGCAGCTTA TTGCGCGCCACG AA GATTGCGTGC GGG GTGATGGGTAGTGCCGACGAATTTAACG TTGGATGTAGTTCAAGTCCGC GC GATGCAATG