File: 21-MI-1_1813F_premixed.ab1

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

Signal G:1451 A:2628 C:2877 T:3115

macrogen

Sample: 21-MI-1_1813F_premixed

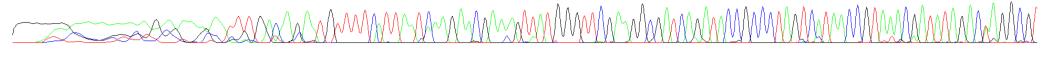
Lane: 71

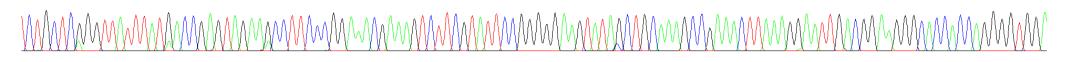
Base spacing: 15.678979

867 bases in 16860 scans

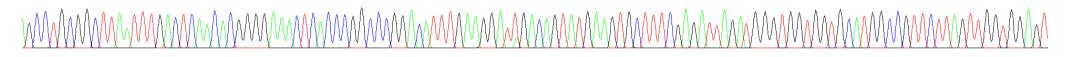
Page 1 of 2

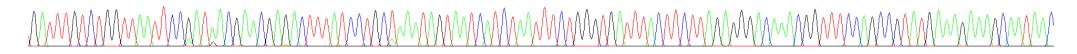
10 20 30 40 50 60 70 80 90 100 110 120 GOGGG AGG C A GCATAC T GC GTA GGA T TG GAC A GAAT GT T T CAT T AAT CAA GAAC GAAAAGT TAT G G GT C GAA GGC GAT CA GAT AC C GC CCT AG T CAT AAC C GT AAAC GAT AT A GAC AA GGC GT CAT AAC C GT AAAC GAT AT A GAC AA GGC GT CAT AAC C GT AAAC GAT AT A GAC AA GGC GT CAT AAC C GT CAT AAC C

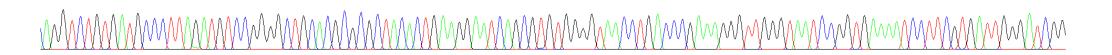




260 270 280 290 300 310 320 330 340 350 360 370 GCCT GCGGCTTAATTT GACT CAACAC GGGGGAAACT CACCC GGCC CGGACATT T GAAGGATT GACAGT AT GAAGT GCTTT C GA GATTAGAT G GGT GGT GGT GGT GCC GTT CTT AGTT GGT GGAGT







File: 21-MI-1_1813F_premixed.ab1 Run Ended: 2022/6/9 16:39:58 Signal G:1451 A:2628 C:2877 T:3115 Sample: 21-MI-1_1813F_premixed Base spacing: 15.678979 Page 2 of 2 *Lane: 71* 867 bases in 16860 scans 630 640 650 660 670 680 690 700 710 720 730 740 750 GACTTGCAATTATTTCCCTTGAACGAGGAATCCCTAGTAAGTGCAGGTCATCAGCTTGCGCTGATTACGTCCCTTGTACACACCGCCCGTCGCTGCCCGGAACTGAACCATTTCGAG 750 760 770 780 790 800 810 820 830 840 850 860 AAATGTGGGGGACTGACGACTTGACGGGCTTCTTGTTGACGGGAAACCGCCTTAATCGCAGTGGTTTGAACCGGGCAAAAGTCTACAC GGGGGGA GAAAAAAA G AAAAAAA