

Run Ended: 2023/4/4 0:17:12

Signal G:1518 A:2154 C:2620 T:2220

macrogen

Sample: 22CCRAA000_D07_premix

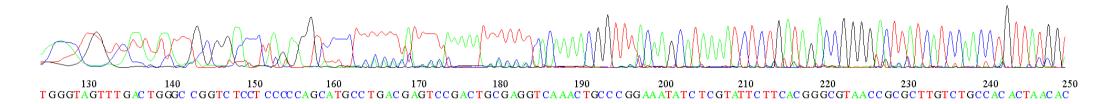
Lane: 57

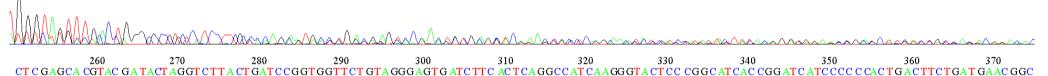
Base spacing: 14.890465

1303 bases in 16300 scans

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File: 22CCRAA000_D07_premix.ab1

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1303 bases in 16300 scans

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 $630 \qquad 640 \qquad 650 \qquad 660 \qquad 670 \qquad 680 \qquad 690 \qquad 700 \qquad 710 \qquad 720 \qquad 730 \qquad 740 \qquad 750$ A GAGAT AACCA CCT GGTG CC T C GCCA C CG GGAG CTT GCCTTCATT AT GGTTACATAC A T GA GTT CGTG AA T GCCCCCC ACC G A A G GTT AAAAAA GA CAAAATTTGGGAAA AA TTGGGTGAG A T GAA T A A

760 770 780 790 800 810 820 830 840 850 860 870 TTGTTGTGG GGC GTT ACC CTC ACCGAAA C AATCT GC CGGT GTAATCAAATGTT GTGTTTTT C C CG T CC AA TCCCCCC A GGCAAAATT GTAT GGT T T A A A GAA AA C C C C C CA GAA GTT A TACCCC

880 890 900 910 920 930 940 950 960 970 980 990 CAAACCAAGAACCTTCG AATTTTT CA A GGGGGATTTT GGGGGGATCT AG AA T CTC GTG CAATTTTT C ACT GGAAGGT ACTTTT CCCCCCTTAAAAT G TTTGC G CC GTTA AT CTTCCC AA

1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120 CATATTC T GCCTCTGAT AT GGCCCC TGG CCTTTA AG AACC GTAGAA GA C ATTA AGCTTC G TCCCA CTA CAA GTTA CTTCTG GA AT AT GGCA GAT GT CCCAC C CTCT C A AT TTTT A CACC TCTCCCCC

1130 1140 1150 1160 1170 1180 1190 1200 1210 1220 1230 GGTG AGA AG GTG ACCT AACTCTGTCCTTA CGA AC A TT CTGA AA C CTCACGCT TT GC G AACTC TA GT TT CAAT AG TAA AAG AA G C C T CTAT CT CAT TGG GA ACT C GT T