

File: 22CCRAA000_H10_premix.ab1

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

Signal G:773 A:816 C:1864 T:1952

Sample: 22CCRAA000_H10_premix

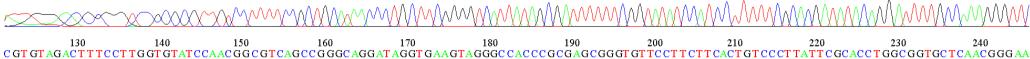
Lane: 66

Base spacing: 15.219386

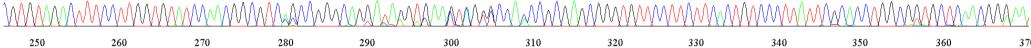
1841 bases in 22576 scans

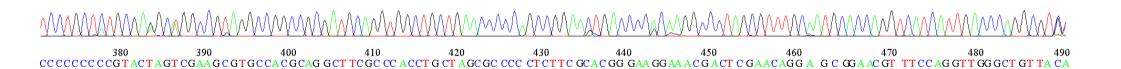
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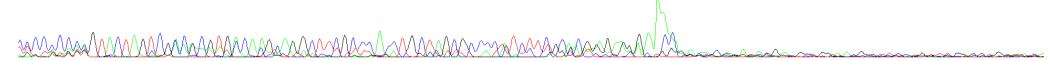












500 510 520 530 540 550 560 570 580 590 600 610 G G C CAC AG GA A G G G A A A A G G G A A C A T C C G A G C G G C G C A G C G A G C C G T C G A G A A C C G T C G A G A A C T G T G C G T G T A T G C G A G C G G C G A G A G A T T T A C G A G C A G

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GATCAGAAA CGCCC CTGCTGAA GATGACAT GGGGGATG GTGTTAGACGT CG GT CAT CGAAACGTA AGGT GA AACTAGT GA TAGTTAGACC GAAAC GTGCAA GGAAGAGGCTGC GGGTG GGAG

GA GCG AT GA GGA T AACA GA C AT GT TAC T GACE TAT GGT AGTG A GA ATGGAAT AGGA GT ACT AAA G A GA GA GGT GAGAT GGAA GAG GT GAGGG G C CA G T GAGCT AGGG A C TGA GAGGA CTG

T GGAAT AGG CGGA C GT GGT AT GT CGA GC AT G GGA T AG GAT GGC T GGA C AGA AACG CT T GAT AGC AGAC GT AC T AT G GCG AATT C AT G AAA G CG GAA C A CGACGT GGACG C GGAT AC GAT G C GA

TGA A GA AG AG G GTAC GG A T AT GA A TG GA A CG GAA C AAA T GGA GAC A G G GGA C T GGA T TG GGA G G GAG A GGAG GCC AGGCT C A GC G TA C C GAT G T G A AA T GGC G AT GCCCT T G A G CG TG T