File: smA_rbcLR.ab1 Sample: smA_rbcLR

Run Ended: 2023-05-31 19:51:33

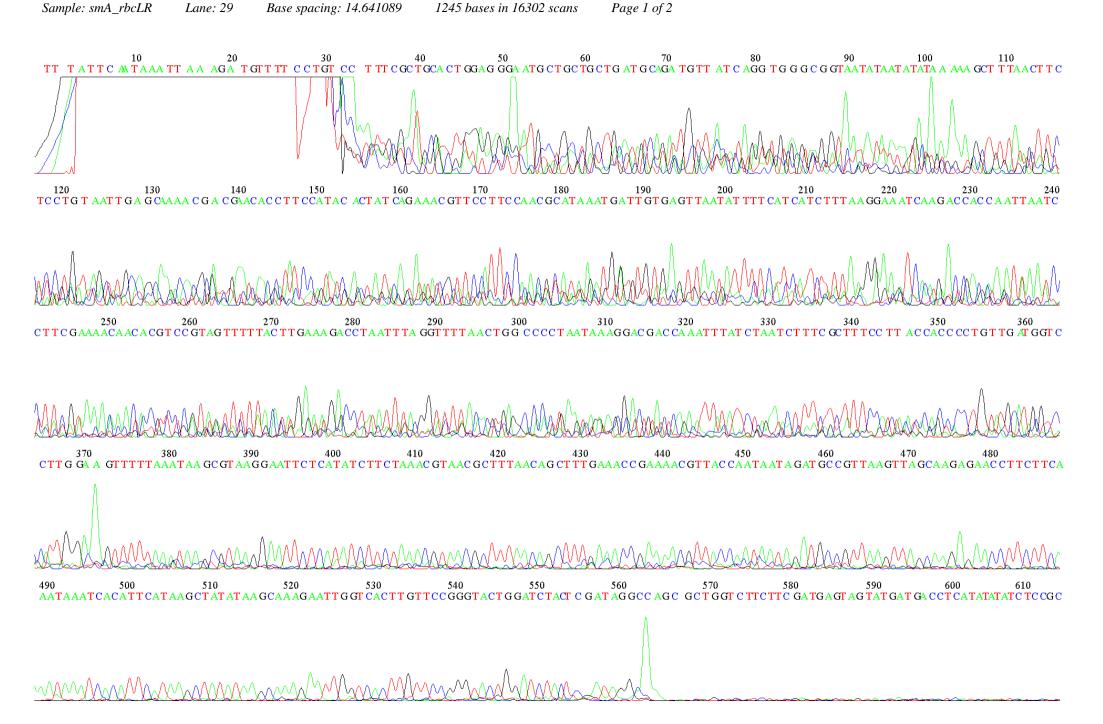
Lane: 29

Signal G:476 A:505 C:549 T:420

1245 bases in 16302 scans

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File: smA_rbcLR.ab1

Run Ended: 2023-05-31 19:51:33

Signal G:476 A:505 C:549 T:420

Sample: smA rbcLR

Lane: 29

Base spacing: 14.641089

1245 bases in 16302 scans

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750 760 770 780 790 800 810 820 830 840 850 860 870 TT AT GAGAT GA A GGG ATGG TTT C T AAT CC A TA C AGGA GGG TA GCAT GGA GTT GTA TT AG CA A A GCCA AC T CC GC AAT CGT CTT AA ACG AT AC ACG AT AC TGTT G GT CCC GCT TG CTC ACTA CCCA CTAC C C

880 890 900 910 920 930 940 950 960 970 980 990 1000 A CTTGC ACTA GG A GC GGTA TTTGCT CT GAT T CCTCGCA T GAT CCCCTG ATA CCA CAG GTT CA C TGTA ATACCT A A GACC GA A A G AT GTT AATACT AT T TCT GC AGTGT TAGTT G ACTTC TT C CG CA

1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120
T CCTTCCCTC A GGCTT CC AG GAT CCCC AGGAT GCCCCGA CC ACCCT CA AAT AC CACC AT GG A CGAAGTC TT AGAGGG GG TTC C CAAGT A GGT C CAC GATAC AT TC C C C C C G T TGAT T C C CCC