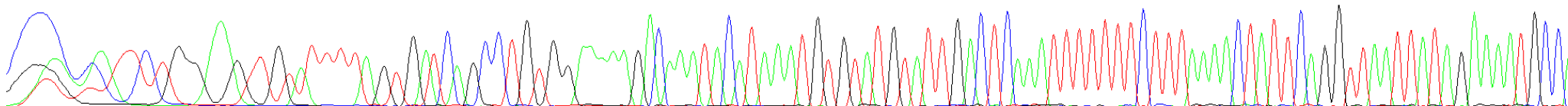
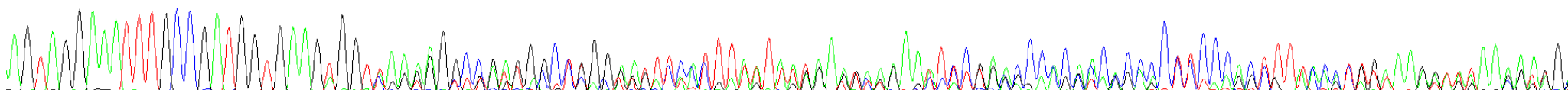


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Sample: 1E59ZAE461_premix Lane: 39 Base spacing: 14.247064 975 bases in 16301 scans Page 1 of 2

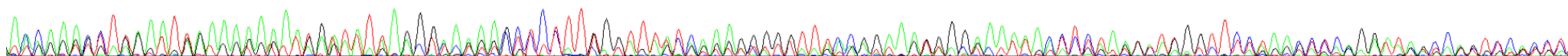
10 20 30 40 50 60 70 80 90 100 110 120
G C T A T A T T C T G G A G T G T A T T T T A G T G A T C A G C C T G T G G A A A A A G A C A A A T A C A T A A A T G T G T A T G T A T T G A C T C A A A T T T T T T C T T T A A A A C T A T T C A G G T T A A T T A T A G A A A T G C C A



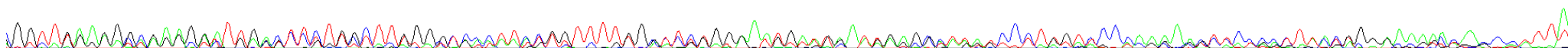
130 140 150 160 170 180 190 200 210 220 230 240
A G T A G G A A A T T T G C C G A T G G T G A A G T G G T T A A G A G G C C G A G G G C C G G G A A T T T C A T T T A T T A A T A A A G A A A A A T T C T A C C C C A C A A C A C C C C C T C C C A C T T T T A C A T G G A A A A G T A T A A G A A T G A



250 260 270 280 290 300 310 320 330 340 350 360 370
A C C A A A C C T G G A A T G G A A A A A A A T G T T A T T A A G T A A A A A G C T C C T T G G G T T T A T C A A A G G G G T T A C C G G A A A T G G G A A A A T A A C T T T A A T A T T G G T T C C A A C C C C A G G T T A G C C A G T C C A C C T C



380 390 400 410 420 430 440 450 460 470 480 490
G G T T G A A G G G C G A A G G A T T G A C T T C T G G C T T G G G G T G A C T T C T G T T T T G G C T T C T G G G A T T G A A G A A T T G A G C C A A A T T C C T T G T C C C A A G A A C T T C C C C T T T A C G G A C A G C C A G C A A T T A



500 510 520 530 540 550 560 570 580 590 600
G C G A T C C G C C C C T G A C G A G C A T C A C A A G A T C G A C G C T C A A G T C A T A G G T G C G A A C C A A C A G G A C T A T A A A G A T A C C A G A C G T T T C C C T G G A A C T C T C G T G C G C T C T C C T G T T



File: 1E59ZAE461.ab1 Run Ended: 2020/2/13 6:50:5 Signal G:620 A:1735 C:1400 T:2509

Sample: 1E59ZAE461_premit Lane: 39 Base spacing: 14.247064 975 bases in 16301 scans Page 2 of 2

610 620 630 640 650 660 670 680 690 700 710 720
CCG AAC TGCCGCTTACC G ATACC TGTCGCCC AT CT CATT C AG AAGC GTGGC GC TT CTCATAG CTCAG CTGTAGGTATC TCAG TT CG G CG AG GT CGT TC GCT CC AG CT G G CTGTG

730 740 750 760 770 780 790 800 810 820 830
TGC AC GAGA CAGT TGT TCAGC C GACCGCTGCGC CTTAT C C GTA ACTATC G TATTG AGTC G AC CTG GTTA GACACG CTT T C G CACTG C ATCAGA TC TG GAAAC G G ATTAT CAG AG

840 850 860 870 880 890 900 910 920 930 940
CG T GT ATGTA TGCAG TGCTA CAGAG TCTTAGA AT G G GATCTA CTA C GCTACACT G A GTAGCA G G ACT G A C TCTCGGCTC TGCTGAAGC G T A C TTCT G AAA A GAGT TC

950 960 970
G T GC GC TG AT CG GC A A CA A T A C GCTT G CAGC