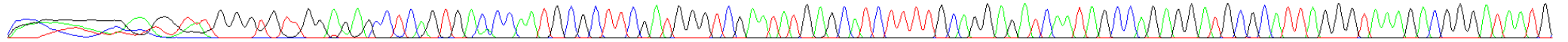
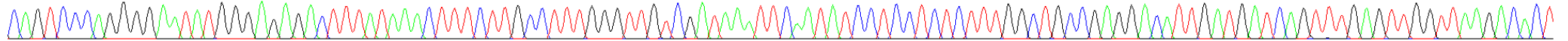


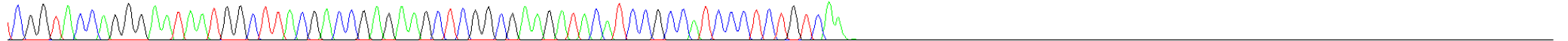
10 20 30 40 50 60 70 80 90 100 110 120 130
C CC GTAGGTGTA TG TATG GGTGTTGGAGAGCTCA GTGACACCAATG C GCTTCGATGGGTTCGAATATGAGCATCTTTTGCAGGAGATCAATAGCCAGAGGATGC GCATTAGGGTAAAGACGGGATAATG



140 150 160 170 180 190 200 210 220 230 240 250
CAGTCCCAGGGGAAATATGGGAGAGACTTTATAAACTTTCTTGCCCTTTGGGTTGTCGATAAATTCAATATCCTCCTCTCTTTGGCTGCCGAGGACATTGATGATCAGTTTGAGTTGTTAAGACACT



260 270 280 290 300 310 320 330 340 350 360 370 380
CGGTACCAAGGGAATAATGGCTTACGACCGAGAAAGCTCGGCGAAGATACATCCGCCATCCCTCTGTCAAAGAGGTGCGGCATTGAGGATCACACATCGTCATCGCATTTTCTTTGTTTTCACGA



390 400 410 420 430 440 450 460 470 480 490 500
CA C CTGA A CCAGG CATATCATGT CACA GCT A TATTTGTGTG TGGGC TGC AATACGTG ATGCC CAC TTGCC ATTCAC AACCATG TAGAA AAAGTAATGT TGTTTGTCTGCGCATGTCTAT A

510 520 530 540 550 560 570 580 590 600 610 620 630 640
ACACGTGGGGCTGA TTTGTTGAATT C GAA TAGAAA AATGGTGGTCC T G TTTAAACGTTAG TGACG C GTTACATTATGCTAA CTTGATACTAAATAATATG TCA GCGG GGTCTGAGGCTAAGATCATATATCGAC

File: ORF3_ORF_R.ab1 Run Ended: 2025/5/10 3:56:54 Signal G:426 A:420 C:343 T:437

Sample: ORF3_Plate1_G10_ORF_R Lane: 68 Base spacing: 15.274415 1576 bases in 16136 scans Page 2 of 2

650 660 670 680 690 700 710 720 730 740 750 760 770 780 790
A CT A CATCAAATAATCTAGTCTCCGTC TTT CTGGT GACGCTA TTCTACCAACATGCAAAATGACGACCTTCCAGCC GGTCAA GC ACATATTGAA TC GATCAAGGTC TAAGAACTGTTGAAATGTCGG TGCTTCTAT CATTGTGACG

800 810 820 830 840 850 860 870 880 890 900 910 920 930 940 950 960
ACATGTC TGGAATCAGGTCCGTCACCTCGAATCTACCATCCGATG GGCACGGTGTCTTCAACTATCTGAGTATCCCCAGTCA TTCTGTGACCTTCACGCGTAGGCGCCATGAGCA TGTCTATCGGACCGAGAACTAGTCCCTCATG AA

970 980 990 1000 1010 1020 1030 1040 1050 1060 1070 1080 1090 1100 1110 1120 1130
TAA A GACTATCAACGCACGCACTTGAA TCATGTTCTAGATCTGT GCGTGATCGTC TCCGCA GG CATA TTGTT GTATTAGTACCCCTTGCTATT AAGGTG TGAATGAGGTACGT CGTTGGGTCTTCGATAATTTA AAGCCGTATTT C T CACGT

1140 1150 1160 1170 1180 1190 1200 1210 1220 1230 1240 1250 1260 1270 1280 1290 1300
GAAATCGT TG GCTTTGTGT GTCATGTCCGTGATCTCA GACTGACACTCTCGGTCTGGGTG TTGTCGCTCCGAGA CCATGGACGCATTAAATTGATTAACCTTCACACGAGATTCATT CAGTCTA TTATTATAATGTGATATGATTCATTCCCTCG

1310 1320 1330 1340 1350 1360 1370 1380 1390 1400 1410 1420 1430 1440 1450 1460 1470
CATCTGTATGAGTGCATCTAGCGTACGTTGTTCTT CTTG GCGGT TATGGTTCTATAATATGATGTTTG GGTCCGATGTGTATATTAGTTTGTGTACACAGGT ACATGTCATA CTCTCGTCTATAAAGAGTTAATGGATCACTCA C C C TGGCG