



630 640 650 660 670 680 690 700 710 720 730 740
AACTACCAGATC CC AAA AA GG AGC GCTCCC GGACTG AGCG ACAACT CAAA GA TA AA ACCACCAC CTGCACACAG AA CA CGTAC AATCAA ATAA CAGACATTA GACATT C GCA GGAGAC CCC

750 760 770 780 790 800 810 820 830 840 850 860
ACGAGACG AACGT TC GACAGT CACCATA GAAA CGTC GCAC AACAA GATT ATT AGCCTACT CCTTAACAGGAT GGCACGGA GAGCAGCCC ATAGCACAAAAAGCTTCGTGATG GAGCAA CGA

870 880 890 900 910 920 930 940 950 960 970 980
CTATTTT TTA GATAGGAC TAGC AGGAT TAA G TGT TTATCAAT AACAA GGGTAC CGTGT TGC GATTCTT AGG AAC CAG GGGAA CCA TAA CCAACG TTCTTTT TGA AAAAAA ATAAAAAT

990 1000 1010 1020 1030 1040 1050 1060 1070 1080
TGT CCC T CATT T G AGGTGG C T T T T T T T A A A A G G G A G A T A A A A G G G A A A T T T A A A A A A A A A T A A C A G G A G A C C T A A C C T A A T A G T A C T T A T C T A A

1090 1100 1110 1120 1130 1140 1150 1160 1170 1180 1190
GGAGGAAA CTTT T ACCC CCC CCA G CG T GGATGTG C GGAG AA AAA AAAAA AAA AA AAAA GTT A TT AT ATT TT TTTTA TATTA TGTTG ATCA CAAGAAA A A AAA ATAA CA T