



630 640 650 660 670 680 690 700 710 720 730 740 750  
CA AATAC GAAGCCGTCCCAAATGAATCCCCCAAAGCATCACGC CCGTGTTCATCCCCCTTCAGTC A AATGGACAATTTCCTCAA GCCGCTTGGACCTTCTGCATTGCCGTGCGCCATCAGCGGTGG TGCAG

760 770 780 790 800 810 820 830 840 850 860 870  
TCCGAGCACGACAACCA A CCCC TTGAACATCGACGCCACCGCAGATCGAAGACA GCGCACTGAATGCCTGCCGACGATGCCAGCCAGGTGTGCTATCCGTGAGAGCACACGCGCTCCACTGGGCT

880 890 900 910 920 930 940 950 960 970 980 990 1000  
GGCAC TGA C G C G C G CA GAC C C G C A T C GGTTACC TCCATCACCGCCTGCA GAGTGATGGACGC TCCTGCCCTACCACACA CGCCGGCTATGA A CCCCCCCCACG C G C TACA CGACCATCAAT

1010 1020 1030 1040 1050 1060 1070 1080  
C ACAG AAT CGCTG CCCCCCAACAGAAAC A T GC G T TGT TCATA TAC GATTAGTGA TGC GAATCGA A A AAC AC AC CGGATAGT T C