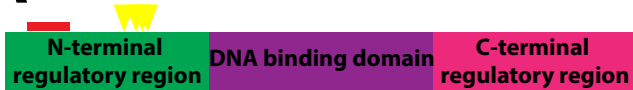


A



B

1	AAGAGCCGAA	TGCACATAAG	GTCGCCAGCC	CACCCTCCGG	ACCCGCATAC	CCCGATGATG
61	TCCTGGACTA	TGGCCTCAAG	CCATACAGCC	CCCTTGCTAG	TCTCTCTGGC	GAGCCCCCG
121	GCCGATTCGG	AGAGCCGGAT	AGGGTAGGGC	CGCAGAAGTT	TCTGAGCGCG	GCCAAGCCAG
181	CAGGGGCCTC	GGGCCTGAGC	CCTCGGATCG	AGATCACTCC	GTCCCACGAA	CTGATCCAGG
241	CAGTGGGGCC	CCTCCGCATG	AGAGACGCGG	GCCTCCTGGT	GGAGCAGCCG	CCCCTGGCCG
301	GGGTGGCCGC	CAGCCCGAGG	TTCACCTGCT	CCGTGCCCGG	CTTCGAGGGC	TACCGCGAGC
361	CGCTTTGCTT	GAGCCCCGCT	AGCAGCGGCT	CCTCTGCCAG	CTTCATTTCT	GACACCTTCT
421	CCCCCTACAC	CTCGCCCTGC	GTCTCGCCCA	ATAACGGCGG	GCCCGACGAC	CTGTGTCCGC
481	AGTTTCAAAA	CATCCCTGCT	CATTATTCCC	CCAGAACCTC	GCCAATAATG	TCACCTCGAA
541	CCAGCCTCGC	CGAGGACAGC	TGCCTGGGCC	GCCACTCGCC	CGTGCCCCGT	CCGGCCTCCC
601	GCTCCTCATC	GCCTGGTGCC	AAGCGGAGGC	ATTCGTGCGC	CGAGGCCTTG	GTTGCCCTGC
661	CGCCCGGAGC	CTCACCCAG	CGCTCCCGGA	GCCCCCTCGC	GCAGCCCTCA	TCTCACGTGG
721	CACCCAGGA	CCACGGCTCC	CCGGCTGGGT	ACCCCCCTGT	GGCTGGCTCT	GCCGTGATCA
781	TGGATGCCCT	GAACAGCCTC	GCCACGGACT	CGCCTTGTGG	GATCCCCCCC	AAGATGTGGA
841	AGACCAGCCC	TGACCCCTCG	CCGGTGTCTG	CCGCCCCATC	CAAGGCCGGC	CTGCCTCGCC
901	ACATCTACCC	GGCCGTGGAG	TTCCTGGGGC	CCTGCGAGCA	GGGCGAGAGG	AGAAACTCGG
961	CTCCAGAATC	CATCCTGCTG	GTTCCGCCCA	CTTGGCCCAA	GCCGCTGGTG	CCTGCCATTC
1021	CCATCTGCAG					