

Supplementary Figures

Figure S1

1 GAGAGATTTC TGGGCAAGAA TTAGTTCTTC TGTTTTTGAA ATTAATTAAC TGAGACAGGA
.....
541 CATTGTGTGC TACGGTAAAA ACAACCCAAG ACAACTCTTC TGAAGCTGAA TCTGCCAGTC
601 ACTCTAAATC TGCAC TATTG CAATACGACT TCATTAATTA CTGCTCAAAC ACTGGCCTAG
TATA Box
.....
901 GTGGCGCTGC CGTCCCAGCC TTGGTCCACG GCCGCCCGCC ACTTCCCTGC CTCTGCGGGA
961 GCCCCCAGC TTGGTCTCCC GCAGGCCGGC GGTGGGGCAC GAGCCCGGTA GGGGGCGCCG
1021 CCGGCCAGG CGCCGCTGCT CCCTCCGCC Sp1 ACCCGAGGC AP2 GGCTGCCGCT AGCCGAGGCC
1081 GGC GCGAGTG CACTGCGAGG CGGGCCGAG GGA GGGAGGC GCGAAGAGGG CGCGAGGTA
E2F (-22)
↪ TSS (-1)
1141 GCAGCCAGAG GGAGCCGCCA GCCCTGCCTG CGGATCCCCG CCGGGCGCAT GGGGCGCTTC
1201 GAGCCGGGAC TGCGTGC GGG CCCCAGAGCC AGTTTCCTGC TGTCGTAAGA AGCCGCGCCA
1261 GGACGCCCGC CGGACCCCGA GCTGCTGGGA GGATG (Translational Start Site)