

**SUPPLEMENTARY TABLE 1****Supplementary Table 1. Sequence of primers utilized**

Amplified Region	FORWARD (5'-3')	REVERSE (5'- 3')
+2 to +76	CGGGGGTCATCTGGGAATAG	GCATTATCCACAGCCCCGT
+32 to +136	GAGCCGCACCAGATCTTCAG	CTGGGACTCAAGTCACGCCTAC
+773 to + 902	CAGCGATGGCACAGAATCGG	TAGCAAAATGTGGGGGTCTGC
+981 to +1074	G TTCACGCACGACAAAGC	GAACAGAGGGTTTGCCGAAGG
+1172 to +1289	CTCGAGACTTGAGGTGAACCC	GGAGTCGAAACTCGCCAGTCT
+1412 to +1522*	ACAGGAAAGTGGCCGCGA	CAACTGGATGGGAATCGGC
ER non responsive region	GCCACCAGCCTGCTTTCTGT	CGTGGATGGGTCCGAGAAC
*Primers amplifying both ERE half sites		

**Sequence of primers utilized.** The table shows the primer sequences used in the ChIP experiment. The primer sequences are amplifying the half ERE sites located upstream of Transcription Start site (TSS).