

Gene	Region	Primers		E-Box position	c-Myc M00322 hits		EGR-1 M00243 hits	
		Sequence	Position		Sequence (strand)	Position	Sequence (strand)	Position
RAP2B	Promoter	CCACGTGCTTCTCGCAGAG GAGCGAGGAAAGCGCTGTAG	-675 -581	-674	GCCACGCGCG (+)	-296	CTGCGGAGGTGG (-)	-250
	Control	AGTCCGGGAGAACCATATGAAG GGCAGAAAAGGAACCCGATC	1782 1882					
KPNA3	Promoter	CCAGGTCCGGCTCCATTT ATGACCCTGCCCTCTTCCC	-318 -235	-174	GCCGCGCGCG (+) GCCGCGCGCC (+)	-449 -153	TTGTGGGGGCGG (+)	-499
	Control	CATATTGGCCATGCACCCA AGGATGGATGGTCGTTAAGCA	6647 6731					
POLH	Promoter I	GCATATAAGCATAAGGGTACCAGAGG GCTTCGTTCTTTATGGTCTCTCCT	-877 -774	-752	GCCACGCGCC (+)	-277	CCGCGTGGGCGG (+)	-489
	Promoter II	GCGGGATCTACACGTGCG GTCCCTATGCTGCTTCTCATCTC	-762 -604					
	Control	TCAGTCATGCCTATCCAATGAAGT TCCTTTTGCTCCTATCCAGCAG	3246 3327					
PTPN1	Promoter	GTCGAAATTCTCGATCGCTGA ACCGGCAGCCGGA ACT	-58 25	---	CCCACGCGCC (+)	-225	AAGCGGAGGAGG (+)	-151
	Control	GCTGTGAAGAGAACGACTTCC CACAAAGCAAATGTCTCCTCC	945 1057					

PP	Promoter	GGAAGCAGGTGACTTATCAACG CTGTTCCCTTCGGCTTTGG	-366 -191	-171	GCCACGTGGC (+)	-173	GCGCGGGGGCGT (+)	-674
	Control	GATTATCGGCGCTGACGC CCCGATCCGGGCCTC	616 687		GCCGCGTGCG (+)	53	GTGCGCAGGCGT (+)	17
KHSRP	Promoter I	CTAATCCCCTCGCTGTCTGG AGGAAATCAAAGCGCCAC	-723 -635	-836 -440 -346	GCCACGTGCC (+)	-838	CGACGTGGGCGG (-)	-747
	Promoter II	AACTCGTCAGGAATTGGATTAATTG TGGGCGTTAGGTTTGATAGTCG	-493 -393				CGGCGGGGGCGG (+)	148
	Control	CTGACCAACTCCACTGTTATCCAC AACACGCAACCACCACAG	1789 1885					
MCCC2	Promoter I	CACGTGTTAGATGCGATTCACTC GGGCGTTTTGTCCAATTCTTT	-890 -784	-890 -10	GCCATGTGGG (+)	136	-----	-----
	Promoter II	TGTCAACACAACCAATCCAAG AGCAGGTCTCCCCGGC	-206 -120					
	Control	GAGTCCAGGGTCTTAACTGTTGCT TTTCAACCATAGAACCCACCTAGG	2617 2693					