

Table S3. PCR primer sets and sequences

Quantitative RT-PCR			
Gene name	Accession No	Forward primer	Reverse primer
GAPDH	NM_002046.3	ccatgttcgcatgggtgt	ccaggggtgctaagcagtt
B2M	NM_004048.2	ttctggcctggaggctatc	tcaggaaatttgactttccattc
VEGFA	NM_001033756.1	agtgtgtgccactgagga	ggtgaggtttgatccgcata
MYC	NM_002467.3	tgctccatgaggagacacc	ctttccacagaacaacatcg
C-SRC	NM_198291.1	cgagaaagtgagaccacgaa	ttggcgttgcgaagtca
hTERT	NM_198255.2	gcctcaagaccacgtc	ccacgaactgtcgcagt
MCL1	NM_182763.1	aagccaatgggcaggctct	agaactccacaaacctcct
CCND1	NM_053056.2	gaagatcgtgccacctg	gacctcctctcgcacttct
CCNE1	NM_001238.1	cctcggattattgcaccatc	cctctctatttgcccagctc
BIRC5	NM_001168.2	agaactggccttcttggga	caagtctggctcgttctcagt
XIAP	NM_001167.2	tttgggacatggatatactcagtt	agcactttactttatcaccttcacc
Quantitative PCR of genomic DNA			
Gene name	Gene ID	Forward primer	Reverse primer
ACTB	60	ctgttttggcttgttcag	aggaaaccttccctctcta
B2m	12010	ttggttgagaagcagaaaca	cacacagtcaagttcccaaa
CHIP analysis			
Gene name	Gene ID	Forward primer	Reverse primer
VEGFA	7422	gtccgcacgtaacctcactt	ccaaggttcacagcctgaaa
MYC	4609	agggatcgcgctgagtataa	tgctctcgcctggaatttact
RT-PCR			
Gene name	Accession No	Forward primer	Reverse primer
GAPDH	NM_002046.3	ccatgttcgcatgggtgt	ccaggggtgctaagcagtt
SP1	NM_138473.2	ctatagcaaatgccccaggt	tctgggctgttttctcttc
SP3	NM_001172712.1	ttgcacctgtcccaactgta	tgttgcctcttttccaaga
SP4	NM_003112.3	tgaagagttgcctgttct	tcctggttcattactgcctctt