

S1 Table. Primer sequences used in this study.

Target gene	Forward (5' – 3')	Reverse (5' – 3')
<i>TCF4</i>	GAGCAGCAAGTCCGAGAAAG	ATGCTGAAACCTCTTGCGTC
<i>CTNNB1</i>	CCATCTTCCAGGAGCGAGAT	CAGTGATGGCATGGACTGTG
<i>TNIK</i>	GCTATTGAGATCCGGTCAGT	CAGGCTGCAACATTGAAAGA
<i>c-MYC</i>	AATGAAAAGGCCCCCAAGGTAGTTATCC	GTCGTTTCCGCAACAAGTCCTCTTC
<i>TWIST1</i>	ATTCAGACCCTCAAGCTGGC	GAGAGGGGAGGAAATCGAG
<i>SNAI1</i>	ACCTCCAGACCCACTCAGAT	GCAGAGGACACAGAACCAGA
<i>CDH1</i>	TCCGAAGCTGCTAGTCTGAG	CTCAAGGGAAGGGAGCTGAA
<i>CDH2</i>	CCCACAGCTCCACCATATGA	TTCAGTCATCACCTCCACCA
<i>VIM</i>	CGCCAACTACATCGACAAGG	GGCTTTGTCGTTGGTTAGCT
<i>ITGAV</i>	GCCGTGGATTTCTTCGTG	GAGGACCTGCCCTCCTTC
<i>ITGB3</i>	CGCTAAATTTGAGGAAGAACG	GAAGGTAGACGTGGCCTCTTT
<i>GAPDH</i>	GAGTCAACGGATTTGGTCGT	GATCTCGCTCCTGGAAGATG