

gene name	reference sequence number	forward	reverse
Full length <i>TACR1</i>	NM_001058.3	5'AACCCCATCATCTACTGCTGC3'	5'ATTCCAGCCCCTCATAGTCG3'
Truncated <i>TACR1</i>	NM_015727.2	5'GGGCCACAAGACCATCTACA3'	5'AAGTTAGCTGCAGTCCCAC3'
<i>AKT1</i>	NM_001014431	5'TGGACCACTGTCATCGAACG3'	5'TGCTTCTTGAGGCCGTACAG3'
<i>AKT2</i>	NM_001243027	5'GAAGGCTGGCTCCACAAGC3'	5'GGCCTCTCCTTGTACCAATG3'
<i>CD13</i>	NM_001150	5'CTACCGCAGCGAGTACATGG3'	5'GCTCATCGAAGCATGGGAAG3'
<i>CD133</i>	NM_001145847	5'AAATGGGCCCTTCTGAGG3'	5'CCTGGTGATTGCCACAAAAC3'
<i>AFP</i>	NM_001134	5'TGTGCTGGATTGTCTGCAGG3'	5'GCAGATTCTGTTATTTGTTGAC3
<i>DLK1</i>	NM_001190703	5'GCAACCCCCAAATGGATTTC3'	5'GAGGTCACGCACTGGTCACA3'
<i>EPCAM</i>	NM_002354	5'ATCGTCAATGCCAGTGTACTTCA3'	5'TGAGCCATTCACTTCTGCCTT3'
<i>GEP</i>	NM_2087	5'CTGCCAGTGGGAAGTATGG3'	5'TGGATCAGGTCACACACAGTGTCC3'
<i>KRT19</i>	NM_002276	5'GCCACTACTACAGACCATCCA3'	5'AGCCAGACGGGCATTGTCC3'
<i>CTNNB1</i>	NM_001904.2	5'ACGTCCATGGGTGGGACA3'	5'CTAGGATGTGAAGGGCTCCG3'
<i>AXIN2</i>	NM_004655.3	5'TATCCAGTGATGCGCTGACG3'	5'TGTTTCTTACTGCCACACGAT3'
<i>LGR5</i>	NM_003667.3	5'ACAGCAGTATGAGACCTTCA3'	5'CAGGTCCTCCTCAAAGTCAAGCA3'
<i>NANOG</i>	NM_024865	5'AGAACTCTCCAACATCTGAACCT3'	5'TCGGCCAGTTGTTTTCTGC3'
<i>SOX2</i>	NM_003106.3	5'CCGTTCATCGACGAGGCTAA3'	5'TTCTTCATGAGCGTCTTGTTTTT3'
<i>OCT4</i>	NM_002701.5	5'CACTGCAGCAGATCAGCCA3'	5'GCTTGATCGCTTGCCCTTC3'
<i>KLF4</i>	NM_004235.4	5'ATCTCAAGGCACACCTGCG3'	5'CCTGGTCAGTTTACTGTAGCG3'
<i>TBP</i>	NM_003194.3	5'GCCGAAACGCCGAATAT3'	5'CCGTGTTTCTGTTGCTCTCT3'