

HUGO Gene Nomenclature	Size	Forward Primer Sequence	Reverse Primer Sequence	Probe Oligo Sequence
ACTB	131	TGGTGATGGAGGAGTTAGTAAGT	AACCAATAAAACCTACTCCTCCCTTAA	6FAM-ACCACCACCCAACACACAATAACAAACACA-TAMRA
APC	97	TATTGCGGAGTGCGGGTC	TCGACGAACTCCCGACGA	6FAM-ATTACACAACACTTCTCTCTCCGCTT-TAMRA
HIC1	100	GTTAGCGGTTAGGGCGTC	CCGAACGCCTCCATCGTAT	6FAM-CAACATCGTCTACCCAACACACTCTCCTACG-TAMRA
HLTF	95	CGGCGTTCGGAATTTGTT	AAACGCCTCGACTCCCCTAA	6FAM-AGGAGGCGTATCGAGGCGGTTTCG-TAMRA
hMLH1	87	CGTTATATATCGTTCGTAGTATTTCGTGTTT	CTATCGCCGCCTCATCGT	6FAM-CGCGACGTCAAACGCCACTACG-TAMRA
HPP1	87	GTTATCGTCGTCGTTTTGTT GTC	GACTTCCGAAAAACAAAAATCG	6FAM-CCGAACAACGGACTACTAAACATCCCGCG-TAMRA
p16	162	GGTTATGATGATGGGTAGCGTTC	ACAACACCACCAACGTATCCAA	6FAM-TACGGCGCGGAGTTTAATTGCGTC-TAMRA
RASSF1A	98	GTGTTAACGCGTTGCGTATC	AACCCCGCGAACTAAAAACGA	6FAM-ACCAACTACCGTATAAAAATTACACGCGATA-TAMRA
SFRP1	125	TGTAGTTTTCCGAGT TAGTGT CGCGC	CCTACGATCGAAAACGACGCGAACG	6FAM -TTCGTCGTTTTCGGTTTTTTTGTTCG-TAMRA
SFRP2	126	AAAATAAGTTCGGGTTTCGGC GGTAC	CAATAAACGAACAAAACGCGA ACTACG	6FAM -CGTTTTTCGTTAGTATTTGGTCGCGAGGTTCG-TAMRA
SFRP4	111	GGTGATGTTATCGTTTTTGT ATCGAC	CCTCCCTAACGTAAACTCGA AACG	6FAM-CCTCGACCGCCCCACCCCA-TAMRA
SFRP5	135	AAGATTTGG CGTTGGGCGGGA CGTTC	ACTCCAACCCGAACCTCGCCG TACG	6FAM -CCCTCAACCCCTCCCGCACCCTA-TAMRA
WT1	75	TTCGGGCGGTATTTGGG	CCCGCGACTCCTCCG	6FAM-AGTAAATGGGTTTCGACGTGCGGGAT-TAMRA