

HGNC approved gene symbol	transcript	median depth	% covered >10x	% covered >20x	% covered >30x	% covered >50x
<i>APC</i>	NM_000038.6	207.00	100.00	100.00	99.95	99.76
<i>ATM</i>	NM_000051.3	117.00	100.00	99.97	99.61	97.12
<i>AXIN2</i>	NM_004655.4	99.00	100.00	99.95	99.26	92.01
<i>BAP1</i>	NM_004656.4	99.00	100.00	99.31	96.76	90.13
<i>BARD1</i>	NM_000465.4	178.00	100.00	99.94	98.86	94.65
<i>BLM</i>	NM_000057.4	129.00	99.97	99.45	98.56	95.19
<i>BMPR1A</i>	NM_004329.2	188.00	100.00	100.00	100.00	100.00
<i>BRCA1</i>	NM_007294.4	275.00	100.00	100.00	100.00	99.90
<i>BRCA2</i>	NM_000059.3	191.00	100.00	100.00	99.99	99.76
<i>BRIP1</i>	NM_032043.3	135.00	100.00	100.00	100.00	99.86
<i>CDC73</i>	NM_024529.4	92.00	100.00	99.61	98.17	93.82
<i>CDH1</i>	NM_004360.5	109.00	100.00	100.00	99.74	96.81
<i>CDK4</i>	NM_000075.4	104.00	100.00	98.91	95.24	92.07
<i>CDKN2A</i>	NM_000077.4	170.00	100.00	99.50	97.48	92.36
<i>CHEK2</i>	NM_007194.4	92.00	100.00	99.35	96.74	88.79
<i>DICER1</i>	NM_177438.2	137.00	100.00	100.00	99.91	99.74
<i>FH</i>	NM_000143.3	110.00	98.69	94.82	92.55	91.51
<i>FLCN</i>	NM_144997.7	97.00	100.00	100.00	99.88	95.80
<i>HOXB13</i>	NM_006361.5	102.00	100.00	99.99	99.79	98.47
<i>MAX</i>	NM_002382.5	63.00	100.00	98.33	91.66	67.70
<i>MEN1</i>	NM_130799.2	93.00	100.00	99.97	99.58	97.22
<i>MITF</i>	NM_000248.3	106.00	100.00	100.00	99.99	96.37
<i>MLH1</i>	NM_000249.3	153.00	100.00	99.99	99.91	99.64
<i>MLH3</i>	NM_001040108.1	299.00	100.00	100.00	99.94	99.52
<i>MSH2</i>	NM_000251.3	114.00	100.00	99.93	99.52	98.01
<i>MSH3</i>	NM_002439.5	96.00	99.98	99.86	99.33	94.90
<i>MSH6</i>	NM_000179.2	239.00	100.00	100.00	99.88	99.26
<i>MUTYH</i>	NM_001128425.1	121.00	100.00	100.00	99.82	96.81
<i>NF1</i>	NM_001042492.3	125.00	100.00	99.77	98.75	94.96
<i>NF2</i>	NM_016418.5	99.00	100.00	99.98	99.83	97.57
<i>NSD1</i>	NM_022455.4	160.00	100.00	99.96	99.55	98.36
<i>NTHL1</i>	NM_002528.7	90.00	100.00	99.97	99.48	95.16
<i>PALB2</i>	NM_024675.4	239.00	100.00	100.00	99.94	99.29
<i>PMS2</i>	NM_000535.7	113.00	99.80	99.43	98.58	95.20
<i>PMS2CL</i>	NR_002217	70.50	87.50	78.12	65.62	56.25
<i>POLD1</i>	NM_002691.4	81.00	100.00	99.29	95.60	83.73
<i>POLE</i>	NM_006231.4	107.00	100.00	99.73	98.61	93.21
<i>POT1</i>	NM_015450.3	114.00	100.00	100.00	99.83	96.62

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<i>PRKAR1A</i>	NM_002734.4	104.00	100.00	100.00	100.00	99.99
<i>PTCH1</i>	NM_000264.5	106.00	99.97	99.63	98.62	94.23
<i>PTCH2</i>	NM_003738.5	91.00	100.00	99.61	97.98	94.20
<i>PTEN</i>	NM_000314.8	152.00	100.00	100.00	100.00	99.55
<i>RAD51B</i>	NM_001321814.1	96.00	100.00	100.00	99.47	98.77
<i>RAD51C</i>	NM_058216.3	116.00	100.00	100.00	99.92	98.06
<i>RAD51D</i>	NM_002878.3	71.00	99.51	96.76	90.96	78.13
<i>RB1</i>	NM_000321.2	95.00	99.87	98.11	95.14	89.39
<i>RET</i>	NM_020975.6	90.00	99.98	98.87	95.99	92.86
<i>SDHA</i>	NM_004168.4	96.00	99.00	97.00	96.41	94.74
<i>SDHAF2</i>	NM_017841.2	85.00	100.00	99.10	95.64	92.67
<i>SDHB</i>	NM_003000.3	97.00	100.00	100.00	100.00	97.88
<i>SDHC</i>	NM_003001.3	175.00	100.00	100.00	99.88	98.12
<i>SDHD</i>	NM_003002.4	165.00	100.00	100.00	100.00	100.00
<i>SMAD4</i>	NM_005359.6	132.00	100.00	100.00	99.98	99.94
<i>SMARCA4</i>	NM_001128849.2	92.00	100.00	99.90	98.89	92.48
<i>SMARCB1</i>	NM_003073.5	102.00	100.00	100.00	99.68	98.57
<i>SMARCE1</i>	NM_003079.5	147.00	100.00	100.00	99.97	99.72
<i>SPRED1</i>	NM_152594.3	174.00	100.00	100.00	99.99	99.91
<i>STK11</i>	NM_000455.5	99.00	100.00	100.00	100.00	99.84
<i>SUFU</i>	NM_016169.3	109.00	100.00	100.00	99.68	95.77
<i>TERT</i>	NM_198253.3	84.00	99.76	97.73	95.01	87.45
<i>TGFBR1</i>	NM_004612.4	134.00	93.69	93.56	93.56	93.56
<i>TGFBR2</i>	NM_003242.6	149.00	100.00	99.92	99.31	95.99
<i>TMEM127</i>	NM_017849.3	87.00	100.00	99.37	96.67	88.21
<i>TP53</i>	NM_000546.5	130.00	100.00	100.00	100.00	100.00
<i>TSC1</i>	NM_000368.5	117.00	100.00	100.00	100.00	99.83
<i>TSC2</i>	NM_000548.5	101.00	100.00	99.90	98.44	93.19
<i>VHL</i>	NM_000551.3	64.00	99.90	97.78	89.59	71.47
<i>WT1</i>	NM_024426.6	70.00	99.88	98.27	91.39	74.72