

HGNC approved gene symbol	transcript	median depth	% covered >10x	% covered >20x	% covered >30x	% covered >50x
<i>ABCC9</i>	NM_005691.3	111.00	100.00	99.88	98.95	96.04
<i>AKT1</i>	NM_001014432.1	96.00	100.00	99.95	98.69	92.31
<i>AKT3</i>	NM_005465.7	103.00	100.00	100.00	99.98	97.46
<i>APC2</i>	NM_001351273.1	87.00	96.64	92.14	86.68	75.98
<i>ASXL3</i>	NM_030632.3	206.00	100.00	99.74	99.13	97.48
<i>BRWD3</i>	NM_153252.5	87.00	99.96	99.23	97.07	89.56
<i>BSC12</i>	NM_001122955.3	96.00	99.93	98.42	95.34	89.79
<i>CBS</i>	NM_000071.2	122.00	100.00	100.00	100.00	99.96
<i>CCND2</i>	NM_001759.4	89.00	99.96	97.66	91.86	78.02
<i>CCNK</i>	NM_001099402.2	87.00	91.76	89.49	85.90	76.02
<i>CDKN1C</i>	NM_000076.2	50.00	79.27	66.48	56.73	50.20
<i>CHD8</i>	NM_001170629.2	135.00	100.00	100.00	99.99	99.80
<i>COL2A1</i>	NM_001844.5	83.00	100.00	99.76	97.86	86.86
<i>CYP19A1</i>	NM_000103.4	128.00	100.00	99.17	95.20	90.21
<i>DIS3L2</i>	NM_152383.4	104.00	100.00	99.97	99.17	94.49
<i>DNMT3A</i>	NM_022552.4	82.00	99.86	97.74	91.61	81.41
<i>EED</i>	NM_003797.5	111.00	100.00	100.00	99.88	96.84
<i>EZH2</i>	NM_004456.5	100.00	100.00	99.95	99.50	98.61
<i>FBN1</i>	NM_000138.4	116.00	100.00	99.97	99.83	99.11
<i>FBN2</i>	NM_001999.4	111.00	100.00	99.97	99.80	98.72
<i>FGFR3</i>	NM_000142.4	67.00	99.85	98.50	94.54	78.27
<i>FIBP</i>	NM_004214.5	87.00	100.00	99.20	95.42	90.34
<i>GNAS</i>	NM_080425.3	169.00	100.00	99.92	99.53	96.20
<i>GPC3</i>	NM_004484.4	109.00	99.86	98.68	95.69	90.65
<i>GPC4</i>	NM_001448.3	108.00	100.00	99.97	99.70	98.16
<i>GPC5</i>	NM_004466.6	116.00	100.00	99.80	98.81	94.39
<i>GPR101</i>	NM_054021.1	102.00	100.00	99.92	98.19	94.41
<i>HERC1</i>	NM_003922.4	120.00	100.00	100.00	99.92	99.40
<i>HIST1H1E (H1-4)</i>	NM_005321	72.00	100.00	100.00	97.81	89.81
<i>IGF1R</i>	NM_000875.5	101.00	100.00	99.55	98.40	97.20
<i>IGF2</i>	NM_000612.6	63.00	100.00	99.79	98.31	80.76
<i>TMEM94</i>	NM_014738	95.00	99.99	99.68	97.72	88.98
<i>LOX</i>	NM_002317.7	104.00	100.00	100.00	99.75	96.70
<i>MED12</i>	NM_005120.3	82.00	99.94	98.45	94.27	86.12
<i>MTOR</i>	NM_004958.4	94.00	99.99	99.71	98.68	96.20
<i>NFIX</i>	NM_001271043.2	120.00	100.00	99.47	96.92	88.61
<i>NOG</i>	NM_005450.6	126.00	100.00	100.00	99.96	99.57
<i>NPPB</i>	NM_002521.3	55.00	100.00	97.92	89.71	60.03
<i>NPPC</i>	NM_024409.4	70.00	99.83	97.56	87.81	73.05
<i>NPR2</i>	NM_003995.3	129.00	100.00	99.99	99.89	98.88
<i>NPR3</i>	NM_001204375.2	119.00	100.00	99.61	98.49	97.80
<i>NSD1</i>	NM_022455.4	160.00	100.00	99.96	99.55	98.36
<i>PDGFRB</i>	NM_002609.4	102.00	100.00	99.83	98.58	92.95

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<i>PIK3CA</i>	NM_006218.4	135.00	100.00	100.00	99.97	99.87
<i>PIK3R1</i>	NM_181523.3	112.00	100.00	100.00	99.95	99.05
<i>PIK3R2</i>	NM_005027.4	88.00	91.55	89.18	86.84	83.53
<i>PIK3R3</i>	NM_003629.4	113.00	100.00	100.00	99.96	99.90
<i>PLOD1</i>	NM_000302.4	80.00	99.99	99.49	96.38	78.90
<i>PPP2R5B</i>	NM_006244.4	73.00	100.00	99.28	93.54	77.12
<i>PPP2R5C</i>	NM_001352913.2	118.00	98.64	98.29	96.42	94.26
<i>PPP2R5D</i>	NM_006245.4	117.00	100.00	100.00	99.80	97.60
<i>PTCH1</i>	NM_000264.5	106.00	99.97	99.63	98.62	94.23
<i>PTEN</i>	NM_000314.8	152.00	100.00	100.00	100.00	99.55
<i>RNF125</i>	NM_017831.4	106.00	100.00	100.00	99.85	95.44
<i>RNF135</i>	NM_032322.4	84.00	99.56	94.59	86.64	72.31
<i>SETD2</i>	NM_014159.6	280.00	99.91	99.15	98.08	96.96
<i>SHANK3</i>	NM_001372044	74.00	93.46	87.05	78.15	63.69
<i>SHOX</i>	NM_000451.3	0.00	0.00	0.00	0.00	0.00
<i>SOCS2</i>	NM_001270471.2	183.00	100.00	100.00	99.62	92.09
<i>STAT1</i>	NM_007315.3	104.00	99.24	97.55	97.32	97.31
<i>STC2</i>	NM_003714.2	72.00	99.80	97.59	92.13	77.68
<i>SUZ12</i>	NM_015355.4	86.00	98.40	90.23	82.52	75.54
<i>TCF20</i>	NM_005650.3	201.00	100.00	100.00	100.00	99.90
<i>TGFB2</i>	NM_003238.6	130.00	100.00	99.93	98.83	96.85
<i>TGFB3</i>	NM_003239.4	104.00	100.00	99.90	98.55	92.42
<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>TMCO1</i>	NM_019026.4	96.00	100.00	100.00	100.00	97.44
<i>TMLHE</i>	NM_018196.4	81.00	85.86	83.33	81.34	77.44
<i>ZBTB20</i>	NM_001348800.2	171.00	100.00	99.87	99.10	97.44