

HGNC approved gene symbol	Transcript	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x	% covered >50x
<i>ABCC9</i>	NM_005691	601439	94	100	100	98	93
<i>AKT1</i>	NM_001014431	164730	96	100	100	98	97
<i>AKT3</i>	NM_005465	611223	66	100	95	85	64
<i>APC2</i>	NM_005883	612034	80	99	95	89	78
<i>BRWD3</i>	NM_153252	300553	77	99	97	92	82
<i>CBS</i>	NM_000071	613381	77	100	97	93	85
<i>CCND2</i>	NM_001759	123833	83	100	100	100	99
<i>CDKN1C</i>	NM_000076	600856	63	85	77	70	63
<i>CHD8</i>	NM_001170629	610528	100	100	100	99	96
<i>CYP19A1</i>	NM_000103	107910	88	100	100	99	91
<i>DIS3L2</i>	NM_001257282	614184	95	100	100	100	95
<i>DNMT3A</i>	NM_153759	602769	85	100	99	98	90
<i>EED</i>	NM_003797	605984	52	100	94	85	56
<i>EZH2</i>	NM_001203249	601573	96	100	100	97	91
<i>FBN1</i>	NM_000138	134797	90	100	100	99	95
<i>FBN2</i>	NM_001999	612570	90	100	100	99	94
<i>FGFR3</i>	NM_000142	134934	92	100	99	97	92
<i>FIBP</i>	NM_004214	608296	90	100	100	100	96
<i>GNAS</i>	NM_016592	139320	163	100	100	100	100
<i>GPC3</i>	NM_001164618	300037	73	99	93	86	75
<i>GPC4</i>	NM_001448	300168	94	100	99	98	92
<i>GPC5</i>	NM_004466	602446	99	100	98	97	94
<i>GPR101</i>	NM_054021	300393	91	100	100	100	99
<i>HERC1</i>	NM_003922	605109	91	100	100	99	95
<i>HIST1H1E</i>	NM_005321	142220	87	100	100	100	100
<i>IGF1R</i>	NM_000875	147370	74	100	100	97	88
<i>IGF2</i>	NM_000612	147470	70	100	100	100	100
<i>MED12</i>	NM_005120	300188	73	100	98	95	85
<i>MTOR</i>	NM_004958	601231	80	100	99	97	89
<i>NFIX</i>	NM_001271043	164005	117	100	100	99	94
<i>NPPB</i>	NM_002521	600295	103	100	100	100	100
<i>NPPC</i>	NM_024409	600296	89	100	100	97	86
<i>NPR2</i>	NM_003995	108961	95	100	100	99	96
<i>NPR3</i>	NM_001204376	108962	87	100	100	100	100
<i>NSD1</i>	NM_172349	606681	121	100	100	99	96
<i>PDGFRB</i>	NM_002609	173410	87	100	98	95	89
<i>PIK3CA</i>	NM_006218	171834	75	100	100	99	93
<i>PIK3R1</i>	NM_181524	171833	95	100	100	98	94
<i>PIK3R2</i>	NM_005027	603157	73	95	94	92	83
<i>PIK3R3</i>	NM_001114172	606076	90	100	100	99	93
<i>PPP2R5B</i>	NM_006244	601644	74	100	100	98	93

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<i>PPP2R5C</i>	NM_002719	601645	85	99	96	92	84
<i>PPP2R5D</i>	NM_001270476	601646	125	100	100	100	100
<i>PTCH1</i>	NM_001083604	601309	78	100	99	97	93
<i>PTEN</i>	NM_000314	601728	94	100	97	89	78
<i>RNF125</i>	NM_017831	610432	106	100	100	94	93
<i>RNF135</i>	NM_032322	611358	66	100	100	98	87
<i>SETD2</i>	NM_014159	612778	100	100	100	99	94
<i>SHANK3</i>	NM_033517	606230	88	94	88	83	76
<i>STC2</i>	NM_003714	603665	90	100	100	98	85
<i>SUZ12</i>	NM_015355	606245	75	92	89	85	78
<i>TCF20</i>	NM_005650	603107	99	100	100	100	100
<i>TGFBR1</i>	NM_004612	190181	94	92	92	92	92
<i>TGFBR2</i>	NM_001024847	190182	95	100	100	100	100
<i>ZBTB20</i>	NM_001164342	606025	131	100	100	100	100