

Gennaam	Transcript ID (hg38 e110)	Exons	Extras (UTR, etc.)
<i>ABL1</i>	NM_005157.6	1-5	exon 1 - part of 5'UTR
<i>ABL2</i>	NM_005158	2-8	
<i>AFF2</i>	NM_002025.4	4-10	
<i>ALK</i>	NM_004304.5	All Exons (Coding+UTRs)	intron19
<i>BCL11B</i>	NM_138576	1-4	
<i>BCOR</i>	NM_001123385.2	All Exons (Coding+UTRs)	
<i>BCOR</i>	NM_017745.6	8	
<i>BCR</i>	NM_004327	All Exons (Coding+UTRs)	
<i>BRAF</i>	NM_004333.6	All Exons (Coding+UTRs)	
<i>CAMTA1</i>	NM_015215.4	3,8-10	
<i>CBFA2T3</i>	NM_005187	10-12	
<i>CBFB</i>	NM_022845	4,5	
<i>CD28</i>	NM_006139	All Exons (Coding+UTRs)	
<i>CHMP2A</i>	NM_014453.4	3-4	
<i>CIC</i>	NM_015125.5	12,14-21	exon 21 - part of 3'UTR
<i>COL6A3</i>	NM_004369.4	1-15,40-44	
<i>CREBBP</i>	NM_004380	2-7,15-16	
<i>CRLF2</i>	NM_022148	1,3	
<i>CSF1</i>	NM_000757.6	2-7	exon 8,9 - part of 3'UTR
<i>CSF1</i>	NM_172212.3	N/A	exon 9 - part of 3'UTR
<i>CSF1R</i>	NM_005211	9-14	
<i>CTNNB1</i>	NM_001904.4	2,3,4,7,8	
<i>DEK</i>	NM_003472	2,3,9	
<i>DUX4</i>	NM_001293798	1	
<i>EBF1</i>	NM_024007	10-15	
<i>EGFR</i>	NM_005228.5	1,3,7,8,9,12,15-21,24,25	EGFRVIII
<i>EPC1</i>	NM_025209.5	6,7,9,10,11	
<i>EPOR</i>	NM_000121	8	
<i>ERBB2</i>	NM_004448.4	4,5,8,11-27	exon 27 - part of 3'UTR
<i>ERG</i>	NM_004449.4	All Exons (Coding+UTRs)	exon 2,3 - part of 5'UTR
<i>ESR1</i>	NM_000125.4	All Exons (Coding+UTRs)	
<i>ETV5</i>	NM_004454.3	2,3,7-9	exon 2 - part of 5'UTR
<i>ETV6</i>	NM_001987.5	1-7	exon 1 - part of 5'UTR
<i>EWSR1</i>	NM_005243.4	4-14	
<i>FGFR1</i>	NM_023110.3	All Exons (Coding+UTRs)	exon 2 - part of 5'UTR
<i>FGFR2</i>	NM_000141.5	All Exons (Coding+UTRs)	exon 2 - part of 5'UTR
<i>FGFR3</i>	NM_000142.5	All Exons (Coding+UTRs)	intron 17
<i>FIP1L1</i>	NM_030917.4	9-18	
<i>FLT3</i>	NM_004119	14-19	
<i>FN1</i>	NM_212482.4	11-42	
<i>FOS</i>	NM_005252.4	4	
<i>FOSB</i>	NM_006732.3	1,2	exon 1 - part of 5'UTR
<i>FOXO1</i>	NM_002015.4	All Exons (Coding+UTRs)	exon 1 - part of 5'UTR, exon 3 - part of 3'UTR
<i>FOXR2</i>	NM_198451.4	All Exons (Coding+UTRs)	
<i>FRK</i>	NM_002031.3	1-6	exon 1 - part of 5'UTR
<i>FUS</i>	NM_004960.4	All Exons (Coding+UTRs)	
<i>GLI1</i>	NM_005269.3	4-7	
<i>GLIS2</i>	NM_032575	4	
<i>GPI</i>	NM_000175.5	15-16	
<i>GREB1</i>	NM_014668.4	3-10	
<i>GRM1</i>	NM_001278064.2	All Exons (Coding+UTRs)	
<i>HLF</i>	NM_002126	4	
<i>HMGA2</i>	NM_003483.6	All Exons (Coding+UTRs)	exon 5 - part of 3'UTR
<i>HOXA10</i>	NM_018951	2	
<i>HOXA11</i>	NM_005523	2	
<i>HOXA9</i>	NM_152739	2	
<i>IKZF1</i>	NM_006060	1-3, 7-8	
<i>IL2RB</i>	NM_000878.3	2	
<i>JAK2</i>	NM_004972	6-13, 15-20	
<i>JAZF1</i>	NM_175061.4	2-4	

<i>KDM2B</i>	NM_032590.5	10-23	
<i>KMT2A</i>	NM_005933	2-35	6x tiling
<i>KRAS</i>	NM_004985.5	2-4	
<i>LYL1</i>	NM_005583	4	
<i>LYN</i>	NM_002350	8-9	
<i>MAML2</i>	NM_032427.4	1-3	exon 1 - part of 5'UTR
<i>MAML3</i>	NM_018717.5	1-3	exon 1 - part of 5'UTR
<i>MAP3K3</i>	NM_002401.5	All Exons (Coding+UTRs)	
<i>MAP3K8</i>	NM_005204.4	All Exons (Coding+UTRs)	
<i>MDM2</i>	NM_002392.6	All Exons (Coding+UTRs)	
<i>MEAF6</i>	NM_001270875.3	4,5	
<i>MECOM</i>	NM_001105077	1-4	
<i>MEF2D</i>	NM_005920	5-9	
<i>MET</i>	NM_000245.4	2,4,5,6,13-21	intron 13/14 for MET exon 14 splice
<i>MITF</i>	NM_001354604.2	All Exons (Coding+UTRs)	
<i>MLLT10</i>	NM_004641	2-18	
<i>MLLT4</i>	NM_001040000	2	
<i>MN1</i>	NM_002430.3	All Exons (Coding+UTRs)	exon 1 - part of 5'UTR, exon 2 - part of 3'UTR
<i>MRTF</i>	NM_001308142.2	11-13	MKL2 gene name in hg19 --> new gene name MRTF in hg38
<i>MRTFA</i>	NM_020831	5	
<i>MSANTD3</i>	NM_080655.3	All Exons (Coding+UTRs)	exon 1 - part of 5'UTR, exon 3 - part of 3'UTR
<i>MYB</i>	NM_001130173.2	7-16	
<i>MYBL1</i>	NM_001080416.4	8-16	
<i>MYC</i>	NM_002467.6	All Exons (Coding+UTRs)	
<i>MYH11</i>	NM_002474	28-33	
<i>MYOD1</i>	NM_002478.5	1	
<i>NCOA1</i>	NM_147223.3	11-15	intron 14
<i>NCOA2</i>	NM_006540.4	11-16,22	intron 14
<i>NOTCH1</i>	NM_017617.5	26-30	
<i>NOTCH2</i>	NM_024408.4	26-30	
<i>NOTCH3</i>	NM_000435.3	26-30	
<i>NPM1</i>	NM_002520	6	
<i>NR1D1</i>	NM_021724.5	All Exons (Coding+UTRs)	
<i>NR4A2</i>	NM_006186.4	All Exons (Coding+UTRs)	exon 1,2 - part of 5'UTR
<i>NR4A3</i>	NM_006981.4	All Exons (Coding+UTRs)	exon 2,3 - part of 5'UTR
<i>NRG1</i>	NM_001159996.3	1,3	
<i>NRG1</i>	NM_013956.5	10	
<i>NRG1</i>	NM_013959.4	1,3	
<i>NRG1</i>	NM_013962.3	1	
<i>NRG1</i>	NM_013964.5	All Exons (Coding+UTRs)	
<i>NTRK1</i>	NM_002529.4	All Exons (Coding+UTRs)	
<i>NTRK2</i>	NM_006180.6	All Exons (Coding+UTRs)	
<i>NTRK3</i>	NM_001007156.3	15,16	
<i>NTRK3</i>	NM_001012338.3	All Exons (Coding+UTRs)	
<i>NTRK3</i>	NM_002530	14,15	
<i>NUP214</i>	NM_005085	17-19, 29-35	
<i>NUP98</i>	NM_016320	8-17	
<i>NUTM1</i>	NM_175741.3	All Exons (Coding+UTRs)	exon 1 - part of 5'UTR
<i>OGA</i>	NM_012215.5	All Exons (Coding+UTRs)	gene name MGEA5 in hg19 --> new gene name OGA in hg38
<i>P2RY8</i>	NM_178129	1	
<i>PATZ1</i>	NM_014323.3	1,2	exon 1 - part of 5'UTR
<i>PAX3</i>	NM_181457.4	8	
<i>PAX3</i>	NM_181458.4	All Exons (Coding+UTRs)	
<i>PAX3</i>	NM_181459.4	9,10	
<i>PAX5</i>	NM_016734	All Exons (Coding+UTRs)	
<i>PAX7</i>	NM_001135254.2	5-8	
<i>PBX1</i>	NM_002585	3,5	
<i>PCM1</i>	NM_006197	23-37	
<i>PDGFB</i>	NM_002608.4	2,3	
<i>PDGFRA</i>	NM_006206	9-14	
<i>PDGFRB</i>	NM_002609	8-14	

<i>PHF1</i>	NM_024165.3	2,10-12	exon 1,2 - part of 5'UTR
<i>PICALM</i>	NM_007166	16-19	
<i>PLAG1</i>	NM_002655.3	4	exons 1,2,3,4 - part of 5'UTR
<i>PML</i>	NM_002675	2-7	
<i>PPARG</i>	NM_015869.5	1-3	exon 1 - part of 5'UTR
<i>PRDM10</i>	NM_199437.2	10-16	
<i>PRDM16</i>	NM_022114	1	
<i>PRKACA</i>	NM_002730.4	2,3,4	
<i>PRKCA</i>	NM_002737.3	All Exons (Coding+UTRs)	
<i>PRKCB</i>	NM_002738.7	All Exons (Coding+UTRs)	
<i>PRKCD</i>	NM_006254.4	All Exons (Coding+UTRs)	
<i>PRKD1</i>	NM_002742.3	7-18	exon 18 - part of 3'UTR
<i>PRKD2</i>	NM_016457.5	7-18	exon 18 - part of 3'UTR
<i>PRKD3</i>	NM_005813.6	7-19	exon 19 - part of 3'UTR
<i>PTEN</i>	NM_000314.8	5-8	
<i>PTK2B</i>	NM_173174	7	
<i>RAB7A</i>	NM_004637.6	3-4	
<i>RAD51B</i>	NM_133510.4	All Exons (Coding+UTRs)	
<i>RAF1</i>	NM_002880.4	4-12	
<i>RARA</i>	NM_000964	2-5	6x tiling
<i>RARB</i>	NM_000965	1-2	
<i>RARG</i>	NM_000966	1-5	
<i>RASGRF1</i>	NM_001145648.3	7-16	
<i>RBM15</i>	NM_022768	1	
<i>RELA</i>	NM_021975.4	1-6	exon 1 - part of 5'UTR
<i>RET</i>	NM_020975.6	2,4,6-14	
<i>ROS1</i>	NM_002944.3	2,4,7,29-37	
<i>RSPO2</i>	NM_178565.5	2-4	exon 1,2 - part of 5'UTR
<i>RSPO3</i>	NM_032784.5	1-3	exon 1 - part of 5'UTR
<i>RUNX1</i>	NM_001754	2-9	
<i>RUNX1T1</i>	NM_001198679	2-3	
<i>SET</i>	NM_001122821	7	
<i>SRF</i>	NM_003131.4	All Exons (Coding+UTRs)	
<i>SS18</i>	NM_001007559.3	All Exons (Coding+UTRs)	exon 1 - part of 5'UTR
<i>STAT6</i>	NM_003153.5	2-7,15-20	exon 1,2 - part of 5'UTR
<i>STIL</i>	NM_003035	1-2	
<i>SUZ12</i>	NM_015355.4	1-4	exon 1 - part of 5'UTR
<i>TAF15</i>	NM_139215.3	5-10	
<i>TAL1</i>	NM_001290404	2-6	
<i>TCF12</i>	NM_207037.2	4-6	
<i>TCF3</i>	NM_003200	11-18	
<i>TFCP2</i>	NM_005653.5	1-3	exon 1 - part of 5'UTR
<i>TFE3</i>	NM_006521.6	All Exons (Coding+UTRs)	
<i>TFEB</i>	NM_001271944.2	All Exons (Coding+UTRs)	
<i>TFG</i>	NM_006070.6	All Exons (Coding+UTRs)	
<i>THADA</i>	NM_022065.5	24-30,36,37	
<i>TLX1</i>	NM_005521	1	
<i>TLX3</i>	NM_021025	1	
<i>TMPRSS2</i>	NM_001135099.1	1	exon 1 - part of 5'UTR
<i>TMPRSS2</i>	NM_005656.4	2-6	exon 1,2 - part of 5'UTR
<i>TRIM11</i>	NM_145214.3	All Exons (Coding+UTRs)	
<i>TSLP</i>	NM_033035.4	1	
<i>TYK2</i>	NM_003331	18,19,25	
<i>UBTF</i>	NM_014233	17	
<i>USP6</i>	NM_001304284.2	10-12	exon 1 till 10 - part of 5'UTR
<i>USP6</i>	NM_004505.4	2-4	exon 1,2 - part of 5'UTR - also promoter region USP6
<i>VCP</i>	NM_007126.5	14-15	
<i>VGLL2</i>	NM_182645.3	All Exons (Coding+UTRs)	intron 3
<i>VGLL3</i>	NM_016206.4	1-4	exon 1 - part of 5'UTR
<i>WT1</i>	NM_000378	All Exons (Coding+UTRs)	
<i>YAP1</i>	NM_001130145.3	All Exons (Coding+UTRs)	

<i>YWHAE</i>	NM_006761.5	4,5	
<i>ZCCHC7</i>	NM_032226	1-4	
<i>ZFTA</i>	NM_001144936.2	1-5	exon 1 - part of 5'UTR
<i>ZNF362</i>	NM_152493	3-5	
<i>ZNF384</i>	NM_133476	2-8	