

NeoThetis GENES AND GENETIC ALTERATIONS TESTED

GENES TESTED

The following table indicates the genes tested per panel.

<p>NeoThetis PAN-CANCER PLUS 222 genes</p>	<p>ABL1, ABL2, AKT1, AKT2, ALK, ANKRD26, APC, AR, ARAF, ASXL1, ATM, ATRX, B2M, BAP1, BARD1, BCL2, BCL6, BCOR, BCORL1, BCR, BMPR1A, BRAF, BRCA1, BRCA2, BRIP1, CALR, CFBF, CBL, CBLL, CBLC, CCND1, CCND2, CCND3, CCNE1, CD274, CD74, CDC25C, CDH1, CDK12, CDK4, CDK6, CDKN2A, CEBPA, CHEK2, CIC, CSF1R, CSF3R, CTLA4, CTNNA1, CUX1, CXCR4, DCK, DDR2, DDX41, DEK, DHX15, DICER1, DNMT3A, DUSP22, EGFR, EIF1AX, EPCAM, ERBB2, ERBB3, ERBB4, ERCC4, ERG, ESR1, ETNK1, ETV1, ETV4, ETV6, EWSR1, EZH2, FANCA, FBXW7, FGF13, FGF19, FGF2, FGF3, FGFR1, FGFR2, FGFR3, FGFR4, FLT1, FLT3, FLT4, FOXA1, FOXL2, FOXO1, FRS2, FUBP1, GATA1, GATA2, GATA3, GNA11, GNAQ, GNAS, H3F3A, HDAC2, HOXB13, HRAS, IDH1, IDH2, IKZF1, IL3, INHA, INSRR, IRF4, JAK1, JAK2, JAK3, KDM6A, KDR, KEAP1, KIT, KMT2A, KMT2C, KMT2D, KRAS, LUC7L2, MALT1, MAP2K1, MAP2K2, MAP3K1, MDM2, MECOM, MET, MITF, MLH1, MLLT3, MPL, MRE11, MSH2, MSH6, MTOR, MUTYH, MYC, MYCN, MYD88, MYH11, MYOD1, NBN, NCOA3, NF1, NF2, NFE2L2, NOTCH1, NPM1, NRAS, NRG1, NTRK1, NTRK2, NTRK3, NUP214, NUTM1, PALB2, PARP1, PBX1, PDCD1, PDCD1LG2, PDGFRA, PDGFRB, PGR, PHF6, PIK3CA, PIK3CB, PIK3R1, PML, PMS2, POLD1, POLE, PPM1D, PPP2R1A, PTCH1, PTEN, PTPN11, RAD21, RAD51C, RAD51D, RAF1, RARA, RB1, RBBP6, RET, RNF43, ROS1, RPS14, RUNX1, RUNX1T1, SETBP1, SF3B1, SH2B3, SLC29A1, SMAD4, SMARCA4, SMARCB1, SMC1A, SMC3, SMO, SOX10, SPOP, SRSF2, STAG2, STAT3, STAT5B, STK11, SUZ12, TCF3, TCL1A, TERT, TET2, TMPRSS2, TP53, TSC1, TSC2, U2AF1, VEGFA, VHL, WT1, XPO1, ZRSR2</p>
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GENETIC ALTERATIONS TESTED

The following table shows the types of genetic alterations and coding exon coverage for each gene.

<p>ABL1 Full coding exon coverage (NM_005157.6)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>ABL2 Full coding exon coverage (NM_007314.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>AKT1 Full coding exon coverage (NM_001382430.1)</p> <p>Selected non-coding regions covered</p> <p>▲ ●</p>	<p>AKT2 Full coding exon coverage (NM_001626.6)</p> <p>Selected non-coding regions covered</p> <p>▲ ●</p>	<p>ALK Full coding exon coverage (NM_004304.5)</p> <p>Selected non-coding regions covered</p> <p>▲ ● ■</p>	<p>ANKRD26 Full coding exon coverage (NM_014915.3)</p> <p>Selected non-coding regions covered</p> <p>▲</p>
<p>APC Full coding exon coverage (NM_000038.6)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>AR Full coding exon coverage (NM_000044.6)</p> <p>Selected non-coding regions covered</p> <p>▲ ●</p>	<p>ARAF Full coding exon coverage (NM_001654.5)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>ASXL1 Exons 1-2, 4-13 (NM_015338.6)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>ATM Full coding exon coverage (NM_000051.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>ATRX Full coding exon coverage (NM_000489.6)</p> <p>Selected non-coding regions covered</p> <p>▲</p>
<p>B2M Full coding exon coverage (NM_004048.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BAP1 Full coding exon coverage (NM_004656.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BARD1 Full coding exon coverage (NM_000465.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BCL2 Full coding exon coverage (NM_000633.3)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BCL6 Full coding exon coverage (NM_001706.5)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BCOR Full coding exon coverage (NM_001123385.2)</p> <p>Selected non-coding regions covered</p> <p>▲</p>
<p>BCORL1 Exons 2-8, 10-14 (NM_001379451.1)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BCR Exons 1-16, 21-23 (NM_004327.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BMPR1A Exons 4-5, 7-8, 11 (NM_004329.3)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BRAF Exons 1-13, 15, 16, 18 (NM_004333.6) Exons 1-14, 16, 17, 19 (NM_001374258.1)</p> <p>Selected non-coding regions covered</p> <p>▲ ● ■</p>	<p>BRCA1 Full coding exon coverage (NM_007294.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>BRCA2 Full coding exon coverage (NM_000059.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>
<p>BRIP1 Full coding exon coverage (NM_032043.3)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>CALR Full coding exon coverage (NM_004343.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>CBFB Exons 2-6 (NM_022845.3)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>CBL Exons 1-2, 4-16 (NM_005188.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>CBLB Full coding exon coverage (NM_170662.5)</p> <p>Selected non-coding regions covered</p> <p>▲</p>	<p>CBLC Full coding exon coverage (NM_012116.4)</p> <p>Selected non-coding regions covered</p> <p>▲</p>

Selected non-coding regions which are covered by the test are indicated above.

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● Copy Number Alterations

■ Rearrangements

CCND1 Full coding exon coverage <small>(NM_053056.3)</small> <i>Selected non-coding regions covered</i> ▲ ●	CCND2 Full coding exon coverage <small>(NM_001759.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	CCND3 Full coding exon coverage <small>(NM_001760.5)</small> <i>Selected non-coding regions covered</i> ▲ ●	CCNE1 Full coding exon coverage <small>(NM_001238.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	CD274 Full coding exon coverage <small>(NM_014143.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	CD74 Full coding exon coverage <small>(NM_001025159.3)</small> <i>Selected non-coding regions covered</i> ▲ ■
CDC25C Full coding exon coverage <small>(NM_001790.5)</small> ▲	CDH1 Full coding exon coverage <small>(NM_004360.5)</small> <i>Selected non-coding regions covered</i> ▲	CDK12 Full coding exon coverage <small>(NM_016507.4)</small> ▲	CDK4 Full coding exon coverage <small>(NM_000075.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	CDK6 Full coding exon coverage <small>(NM_001145306.2)</small> <i>Selected non-coding regions covered</i> ▲ ●	CDKN2A Full coding exon coverage <small>(NM_000077.5)</small> <small>(NM_058195.4)</small> <i>Selected non-coding regions covered</i> ▲
CEBPA Full coding exon coverage <small>(NM_004364.5)</small> ▲	CHEK2 Exon 2-13 <small>(NM_007194.4)</small> <i>Selected non-coding regions covered</i> ▲	CIC Full coding exon coverage <small>(NM_001386298.1)</small> <i>Selected non-coding regions covered</i> ▲	CSF1R Full coding exon coverage <small>(NM_001288705)</small> ▲	CSF3R Full coding exon coverage <small>(NM_000760.4)</small> <i>Selected non-coding regions covered</i> ▲	CTLA4 Full coding exon coverage <small>(NM_005214.5)</small> ▲
CTNNB1 Exons 3-15 <small>(NM_001904.4)</small> <i>Selected non-coding regions covered</i> ▲	CUX1 Full coding exon coverage <small>(NM_181552.4)</small> Exons 1-18, 20-23 <small>(NM_001913.5)</small> ▲	CXCR4 Full coding exon coverage <small>(NM_003467.3)</small> ▲	DCK Full coding exon coverage <small>(NM_000788.3)</small> ▲	DDR2 Full coding exon coverage <small>(NM_006182.4)</small> <i>Selected non-coding regions covered</i> ▲	DDX41 Full coding exon coverage <small>(NM_016222.4)</small> ▲
DEK Full coding exon coverage <small>(NM_003472.4)</small> ▲	DHX15 Full coding exon coverage <small>(NM_001358.3)</small> ▲	DICER1 Exons 2-26 <small>(NM_177438.3)</small> <i>Selected non-coding regions covered</i> ▲	DNMT3A Full coding exon coverage <small>(NM_022552.5)</small> <i>Selected non-coding regions covered</i> ▲	DUSP22 Full coding exon coverage <small>(NM_001286555.3)</small> <i>Selected non-coding regions covered</i> ▲	EGFR Exons 1-15, 17-24, 27,28 <small>(NM_005228.5)</small> <i>Selected non-coding regions covered</i> ▲ ●
EIF1AX Exons 2, 4-5 <small>(NM_001412.4)</small> ▲	EPCAM Full coding exon coverage <small>(NM_002354.3)</small> ▲	ERBB2 Exons 1-15, 18-24 <small>(NM_004448.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	ERBB3 Full coding exon coverage <small>(NM_001982.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	ERBB4 Full coding exon coverage <small>(NM_005235.3)</small> ▲	ERCC4 Full coding exon coverage <small>(NM_005238.3)</small> ▲
ERG Full coding exon coverage <small>(NM_182918.4)</small> ▲	ESR1 Full coding exon coverage <small>(NM_000125.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	ETNK1 Full coding exon coverage <small>(NM_018638.5)</small> <i>Selected non-coding regions covered</i> ▲	ETV1 Full coding exon coverage <small>(NM_004956.5)</small> <i>Selected non-coding regions covered</i> ▲	ETV4 Full coding exon coverage <small>(NM_001079675.5)</small> ▲	ETV6 Full coding exon coverage <small>(NM_001987.5)</small> ▲
EWSR1 Exons 1-3, 6-10, 12-17 <small>(NM_005243.4)</small> <i>Selected non-coding regions covered</i> ▲	EZH2 Exons 2-4, 6-20 <small>(NM_004456.5)</small> ▲	FANCA Exons 1-2, 4- 14, 16-43 <small>(NM_000135.4)</small> ▲	FBXW7 Full coding exon coverage <small>(NM_001349798.2)</small> <i>Selected non-coding regions covered</i> ▲	FGF13 Exons 1, 3-5 <small>(NM_004114.5)</small> ▲	FGF19 Full coding exon coverage <small>(NM_005117.3)</small> <i>Selected non-coding regions covered</i> ▲ ●
FGF2 Full coding exon coverage <small>(NM_001361665.2)</small> <i>Selected non-coding regions covered</i> ▲ ●	FGF3 Full coding exon coverage <small>(NM_005247.4)</small> <i>Selected non-coding regions covered</i> ▲ ●	FGFR1 Exons 2, 5-9, 12-18 <small>(NM_023110.3)</small> <i>Selected non-coding regions covered</i> ▲ ● ■	FGFR2 Exons 2-9, 11, 13-17 <small>(NM_000141.5)</small> <i>Selected non-coding regions covered</i> ▲ ● ■	FGFR3 Full coding exon coverage <small>(NM_000142.5)</small> <i>Selected non-coding regions covered</i> ▲ ● ■	FGFR4 Full coding exon coverage <small>(NM_213647.3)</small> <i>Selected non-coding regions covered</i> ▲ ●
FLT1 Full coding exon coverage <small>(NM_002019.4)</small> <i>Selected non-coding regions covered</i> ▲	FLT3 Exons 1-15, 17-24 <small>(NM_004119.3)</small> ▲	FLT4 Exons 2-19, 21-30 <small>(NM_182925.5)</small> ▲	FOXA1 Full coding exon coverage <small>(NM_004496.5)</small> ▲	FOXL2 Full coding exon coverage <small>(NM_023067.4)</small> ▲	FOXO1 Full coding exon coverage <small>(NM_002015.4)</small> ▲

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FRS2 Full coding exon coverage (NM_001278356.2) ▲	FUBP1 Exons 1-19 (NM_003902.5) Selected non-coding regions covered ▲	GATA1 Full coding exon coverage (NM_002049.4) ▲	GATA2 Full coding exon coverage (NM_0032638.5) ▲	GATA3 Full coding exon coverage (NM_001002295.2) ▲	GNA11 Exons 2-7 (NM_002067.5) ▲
GNAQ Full coding exon coverage (NM_002072.5) ▲	GNAS Full coding exon coverage (NM_005016.7) Selected non-coding regions covered ▲	H3F3A Full coding exon coverage (NM_002107.7) ▲	HDAC2 Full coding exon coverage (NM_001527.4) ▲	HOXB13 Full coding exon coverage (NM_006361.6) Selected non-coding regions covered ▲	HRAS Full coding exon coverage (NM_005343.4) (NM_176795.5) ▲
IDH1 Full coding exon coverage (NM_005896.4) ▲	IDH2 Exon 2-11 (NM_002168.4) ▲	IKZF1 Full coding exon coverage (NM_006060.4) Selected non-coding regions covered ▲	IL3 Full coding exon coverage (NM_005888.4) ▲	INHA Full coding exon coverage (NM_002191.4) ▲	INSRR Exons 1-6, 8-22 (NM_014215.3) ▲
IRF4 Full coding exon coverage (NM_002460.4) ▲	JAK1 Full coding exon coverage (NM_002227.4) ▲	JAK2 Full coding exon coverage (NM_004972.4) Selected non-coding regions covered ▲ ●	JAK3 Full coding exon coverage (NM_002215.4) ▲	KDM6A Exons 1-13, 15-24, 26-30 (NM_001291415.2) ▲	KDR Full coding exon coverage (NM_002253.4) ▲
KEAP1 Full coding exon coverage (NM_203500.2) ▲	KIT Full coding exon coverage (NM_000222.3) Selected non-coding regions covered ▲ ●	KMT2A Full coding exon coverage (NM_001197104.2) ▲	KMT2C Exons 1-6, 8-14, 16-17, 22-23, 25-59 (NM_170606.3) ▲	KMT2D Full coding exon coverage (NM_003482.4) ▲	KRAS Full coding exon coverage (NM_004985.5) Selected non-coding regions covered ▲ ●
LUC7L2 Full coding exon coverage (NM_016019.5) Selected non-coding regions covered ▲	MALT1 Full coding exon coverage (NM_006785.4) ▲	MAP2K1 Full coding exon coverage (NM_002755.4) ▲	MAP2K2 Exons 1-8, 10, 11 (NM_030662.4) ▲	MAP3K1 Full coding exon coverage (NM_005921.2) ▲	MDM2 Full coding exon coverage (NM_002392.6) Selected non-coding regions covered ▲ ●
MECOM Full coding exon coverage (NM_004991.4) Selected non-coding regions covered ▲	MET Full coding exon coverage (NM_000245.4) Selected non-coding regions covered ▲ ●	MITF Full coding exon coverage (NM_001354604.2) (NM_000248.4) Selected non-coding regions covered ▲	MLH1 Full coding exon coverage (NM_000249.4) Selected non-coding regions covered ▲	MLL3 Exons 1-5, 7-11 (NM_004529.4) ▲	MPL Full coding exon coverage (NM_005373.3) ▲
MRE11 Full coding exon coverage (NM_005591.4) Selected non-coding regions covered ▲	MSH2 Full coding exon coverage (NM_000251.3) Selected non-coding regions covered ▲	MSH6 Full coding exon coverage (NM_000179.3) Selected non-coding regions covered ▲	MTOR Full coding exon coverage (NM_004958.4) ▲	MUTYH Full coding exon coverage (NM_001048174.2) ▲	MYC Full coding exon coverage (NM_002467.6) Selected non-coding regions covered ▲ ●
MYCN Full coding exon coverage (NM_005378.6) Selected non-coding regions covered ▲ ●	MYD88 Full coding exon coverage (NM_002468.5) ▲	MYH11 Full coding exon coverage (NM_002474.3) (NM_001040113.2) ▲	MYOD1 Full coding exon coverage (NM_002478.5) ▲	NBN Full coding exon coverage (NM_002485.5) Selected non-coding regions covered ▲	NCOA3 Full coding exon coverage (NM_181659.3) Selected non-coding regions covered ▲ ●
NF1 Exons 1-37, 40, 44-47, 49, 53 (NM_001042492.3) Selected non-coding regions covered ▲	NF2 Full coding exon coverage (NM_000268.4) Selected non-coding regions covered ▲	NFE2L2 Exons 2-5 (NM_006164.5) ▲	NOTCH1 Exons 2-34 (NM_017617.5) ▲	NPM1 Exons 1-2, 5-7, 11 (NM_002520.7) Selected non-coding regions covered ▲	NRAS Full coding exon coverage (NM_002524.5) ▲ ●

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NRG1 Full coding exon coverage <small>(NM_013964.5)</small> <i>Selected non-coding regions covered</i> 	NTRK1 Full coding exon coverage <small>(NM_002529.4)</small> <i>Selected non-coding regions covered</i> 	NTRK2 Exons 10-11, 13-14 <small>(NM_006180.6)</small> <i>Selected non-coding regions covered</i> 	NTRK3 Exons 14-15 <small>(NM_001012338.3)</small> <i>Selected non-coding regions covered</i> 	NUP214 Full coding exon coverage <small>(NM_005085.4)</small> 	NUTM1 Full coding exon coverage <small>(NM_001284292.2)</small> <i>Selected non-coding regions covered</i>
PALB2 Full coding exon coverage <small>(NM_024675.4)</small> <i>Selected non-coding regions covered</i> 	PARP1 Exon 1-10, 12-23 <small>(NM_001618.4)</small> 	PBX1 Full coding exon coverage <small>(NM_002585.4)</small> 	PDCD1 Full coding exon coverage <small>(NM_005018.3)</small> 	PDCD1LG2 Full coding exon coverage <small>(NM_025239.4)</small> 	PDGFRA Full coding exon coverage <small>(NM_006206.6)</small> <i>Selected non-coding regions covered</i>
PDGFRB Full coding exon coverage <small>(NM_002609.4)</small> <i>Selected non-coding regions covered</i> 	PGR Full coding exon coverage <small>(NM_000926.4)</small> 	PHF6 Full coding exon coverage <small>(NM_001015877.2)</small> 	PIK3CA Exons 2-4, 14-21 <small>(NM_006218.4)</small> <i>Selected non-coding regions covered</i> 	PIK3CB Full coding exon coverage <small>(NM_006219.3)</small> <i>Selected non-coding regions covered</i> 	PIK3R1 Exons 2-9, 11-16 <small>(NM_181523.3)</small> <i>Selected non-coding regions covered</i>
PML Full coding exon coverage <small>(NM_033238.3)</small> <i>Selected non-coding regions covered</i> 	PMS2 Exons 2-4, 6-11, 13 <small>(NM_000535.7)</small> <i>Selected non-coding regions covered</i> 	POLD1 Full coding exon coverage <small>(NM_002691.4)</small> 	POLE Full coding exon coverage <small>(NM_006231.4)</small> <i>Selected non-coding regions covered</i> 	PPM1D Full coding exon coverage <small>(NM_003620.4)</small> 	PPP2R1A Full coding exon coverage <small>(NM_014225.6)</small>
PTCH1 Full coding exon coverage <small>(NM_000264.5)</small> <small>(NM_001083603.3)</small> <i>Selected non-coding regions covered</i> 	PTEN Exons 1-8 <small>(NM_000314.8)</small> <i>Selected non-coding regions covered</i> 	PTPN11 Exons 1-6, 8-9, 11-15 <small>(NM_002834.5)</small> 	RAD21 Full coding exon coverage <small>(NM_006265.3)</small> 	RAD51C Full coding exon coverage <small>(NM_058216.3)</small> <i>Selected non-coding regions covered</i> 	RAD51D Full coding exon coverage <small>(NM_002878.4)</small> <i>Selected non-coding regions covered</i>
RAF1 Full coding exon coverage <small>(NM_002880.4)</small> <i>Selected non-coding regions covered</i> 	RARA Full coding exon coverage <small>(NM_000964.4)</small> <i>Selected non-coding regions covered</i> 	RB1 Exons 1-13, 16-27 <small>(NM_000321.3)</small> <i>Selected non-coding regions covered</i> 	RBBP6 Full coding exon coverage <small>(NM_006910.5)</small> 	RET Full coding exon coverage <small>(NM_020975.6)</small> <i>Selected non-coding regions covered</i> 	RNF43 Full coding exon coverage <small>(NM_017763.6)</small>
ROS1 Exons 32-37 <small>(NM_001378902.1)</small> <i>Selected non-coding regions covered</i> 	RPS14 Exons 3-5 <small>(NM_005617.4)</small> 	RUNX1 Exons 4-9 <small>(NM_001754.5)</small> <i>Selected non-coding regions covered</i> 	RUNX1T1 Exons 2-12 <small>(NM_001198679.3)</small> <i>Selected non-coding regions covered</i> 	SETBP1 Full coding exon coverage <small>(NM_015559.3)</small> <i>Selected non-coding regions covered</i> 	SF3B1 Exons 1-10, 12-21, 23-25 <small>(NM_012433.4)</small> <i>Selected non-coding regions covered</i>
SH2B3 Full coding exon coverage <small>(NM_005475.3)</small> 	SLC29A1 Full coding exon coverage <small>(NM_001372327.1)</small> 	SMAD4 Full coding exon coverage <small>(NM_005359.6)</small> <i>Selected non-coding regions covered</i> 	SMARCA4 Exons 2-10, 12-35 <small>(NM_003072.5)</small> Exons 2-10, 12-36 <small>(NM_001387283.1)</small> 	SMARCB1 Full coding exon coverage <small>(NM_003073.5)</small> <i>Selected non-coding regions covered</i> 	SMC1A Full coding exon coverage <small>(NM_006306.4)</small>
SMC3 Exons 1-25, 29 <small>(NM_005445.4)</small> 	SMO Full coding exon coverage <small>(NM_005631.5)</small> 	SOX10 Full coding exon coverage <small>(NM_006941.4)</small> 	SPOP Full coding exon coverage <small>(NM_001007228.2)</small> 	SRSF2 Full coding exon coverage <small>(NM_001195427.2)</small> 	STAG2 Full coding exon coverage <small>(NM_001042750.2)</small>
STAT3 Full coding exon coverage <small>(NM_139276.3)</small> 	STAT5B Exons 2-6, 9-12, 14-15, 17-19 <small>(NM_012448.4)</small> 	STK11 Full coding exon coverage <small>(NM_000455.5)</small> <i>Selected non-coding regions covered</i> 	SUZ12 Exons 1-2, 4, 7-8, 10-16 <small>(NM_015355.4)</small> 	TCF3 Full coding exon coverage <small>(NM_003200.5)</small> <small>(NM_001136139.4)</small> 	TCL1A Full coding exon coverage <small>(NM_021966.3)</small>

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