



Genome Build GRCh37 (hg19)

<b>Gene</b>	<b>GenBank Accession Number*</b>	<b>Coverage Region**</b>
<i>BIRC3</i>	NM_001165.4	Exons 2–9
<i>BRAF</i>	NM_004333.4	Exons 11–18
<i>CCND1</i>	NM_053056.2	Exons 1–5
<i>CDKN2A</i>	NM_000077.4	Exons 1–3
<i>CRBN</i>	NM_016302.3	Exons 1–11
<i>CUL4A</i>	NM_001008895.2	Exons 2–20
<i>CUL4B</i>	NM_003588.3	Exons 2–22
<i>CXCR4</i>	NM_003467.2	Exons 1–2
<i>DIS3</i>	NM_014953.4	Exons 1–21
<i>EGFR</i>	NM_005228.3	Exons 1–28
<i>TENT5C</i>	NM_017709.3	Exon 2
<i>IDH1</i>	NM_005896.3	Exon 4
<i>IDH2</i>	NM_002168.3	Exon 4
<i>IKZF1</i>	NM_006060.6	Exons 2–8
<i>IKZF3</i>	NM_012481.4	Exons 1–8
<i>KRAS</i>	NM_033360.3	Exons 2–4
<i>MYC</i>	NM_002467.4	Exons 1–3
<i>MYD88</i>	NM_002468.4	Exons 1–5
<i>NRAS</i>	NM_002524.4	Exons 2–4
<i>NSD2</i>	NM_001042424.2	Exons 17–20
<i>PIK3CA</i>	NM_006218.2	Exons 2–11, 14–21
<i>PIM1</i>	NM_001243186.1	Exons 1–6
<i>STAT3</i>	NM_139276.2	Exons 2–24
<i>TP53</i>	NM_000546.4	Exons 4–11
<i>TRAF3</i>	NM_145725.2	Exons 3–12
<i>XBP1</i>	NM_001079539.1	Exons 1–6

\*Reference transcript numbers may have been updated due to database re-versioning. Refer to the patient report for the most updated gene transcript information.

\*\*Default is +/- 5 bps of flanking intron surrounding each exon with some exceptions. *BRAF* exon 18 has -2 bp of flanking intron. *CDKN2A* contains additional coverage for an alternate exon 1 from transcript NM\_058195 (chr9:g.21994138-21994330). *DIS3* contains additional coverage for an alternate exon 2 from transcript NM\_001128226 (chr13:g.73355427-73355494).