



Genome Build GRCh37 (hg19)

Gene	GenBank Accession Number	Exons–Systematic Numbering
<i>CEBPA</i>	NM_004364.4	1
<i>DNMT3A</i>	NM_022552.4	8–23
<i>FLT3</i>	NM_004119.2	14–20
<i>IDH1</i>	NM_005896.3	4
<i>IDH2</i>	NM_002168.3	4
<i>KIT</i>	NM_000222.2	8–11 and 17
<i>KRAS</i>	NM_033360.3	3-Feb
<i>NPM1</i>	NM_002520.6	9–11, intron 10 30bp before exon 11
<i>NRAS</i>	NM_002524.4	2 and 3
<i>RUNX1</i>	NM_001001890.2	1–6, intron 4 c.725–13T>A and intron 5 c.886+1–4del
<i>TP53</i>	NM_000546.4	4–9

Default offset is +/- 10bps around each exon with exception for *RUNX1*, which has certain intron regions of interest. Also, *NPM1* has coverage set to -30bps before exon 11 because of a downstream polymorphic region.

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