

[SETD1B]

Mutation tier

Entity	Tier	Description
DLBCL	1	high-confidence DLBCL gene

Mutation incidence

Entity	source	frequency (%)
DLBCL	GAMBL genomes	5.16
DLBCL	Schmitz cohort	12.77
DLBCL	Reddy cohort	5.81
DLBCL	Chapuy cohort	NA

Mutation pattern

Entity	aSHM	Significant selection	dN/dS (missense)	dN/dS (nonsense)
BL	No	No	0.000	0.000
DLBCL	No	No	2.645	24.278
FL	No	Yes	4.979	128.132

SETD1B Hotspots

Chromosome	Coordinate (hg19)	ref>alt	HGVSp
chr12	122242755	C>A	P38T
chr12	122243045	G>A	E66K
chr12	122243058	A>G	D70G
chr12	122243060	C>T	P71S
chr12	122265950	G>A	A1858T
chr12	122265953	C>T	R1859C

View coding variants in ProteinPaint [hg19](#) or [hg38](#)

View all variants in GenomePaint [hg19](#) or [hg38](#)

