

S6 Table. Co-localisation between c-Myb footprints and histone modifications. FDR-corrected p values , p', and normalised ratios for co-localization between c-Myb footprints and ChIP-Seq peaks for the four histone modifications analysed in K562 cells.

Histone Modification	p'	Normalised ratio
H3K4me3	4.00 x10 ⁻⁴	1.10
H3K4me1	4.00 x10 ⁻⁴	0.81
H3K9ac	0.31	1.01
H3K27me3	0.31	0.80