

CdLS9 interactions

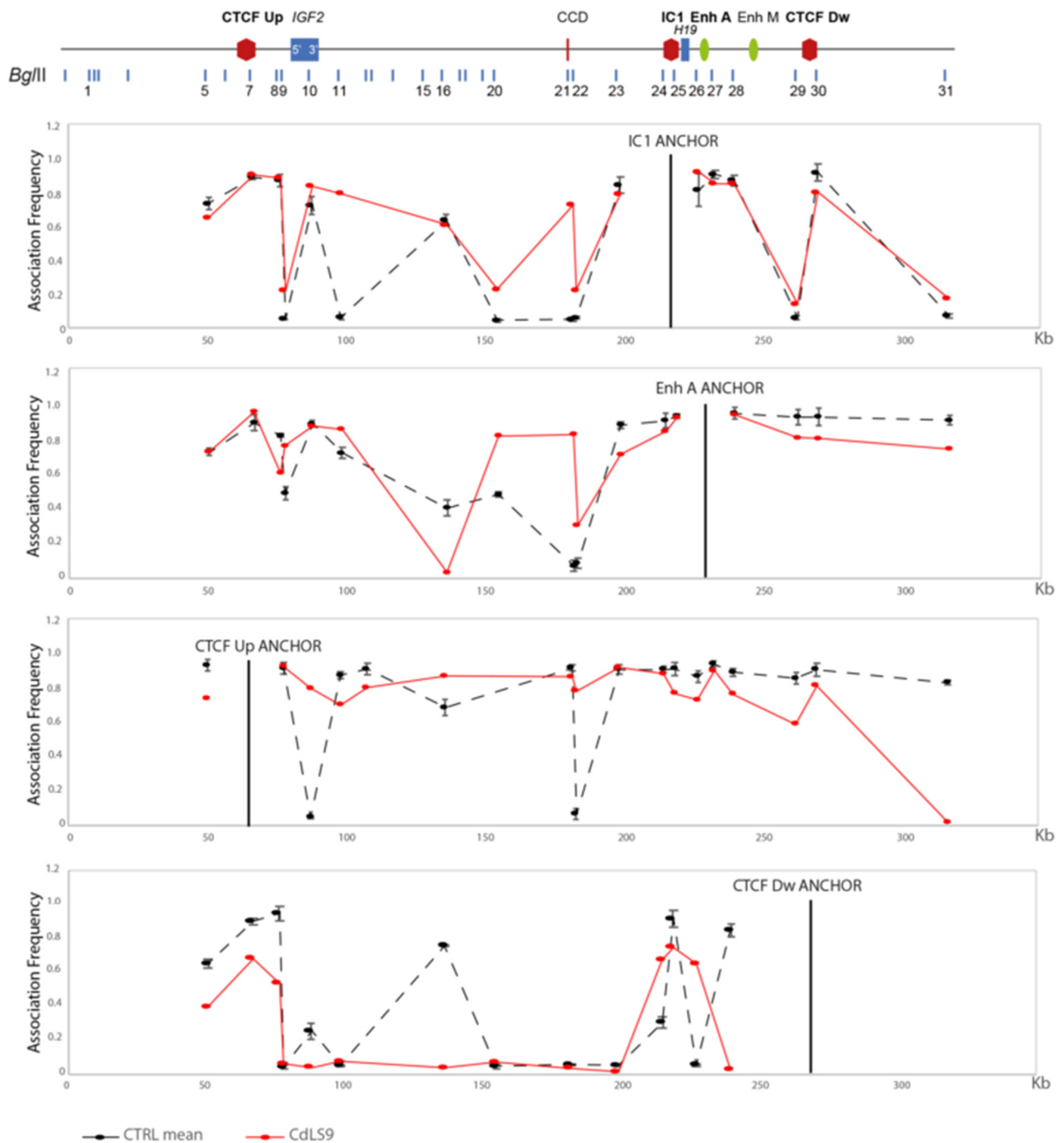


Figure S1. Abnormalities in chromatin architecture at the *IC1* locus of the CdLS9 cell line (*NIPBL* mutation). *IC1* locus looping profiles for the indicated anchors in controls (dotted black) and CdLS9 (red) cell lines. BglII restriction sites are indicated above. Each point in the profile is the mean \pm standard deviation of two independent 3C experiments and indicates the association frequency between the anchor and the fragment on the left of the corresponding BglII restriction site.