



Figure 3 – figure supplement 2 – Comparison of the ends of DNA fragments from DNase I and high-resolution X-ChIP-seq centered over the CTCF motif.

The upper graph shows the average profile, and the heatmaps below show the signal ± 40 bp for each CTCF motif. The leftward shift of the DNase I footprint likely reflects differences in the steric interactions between CTCF and DNase I and that MNase.