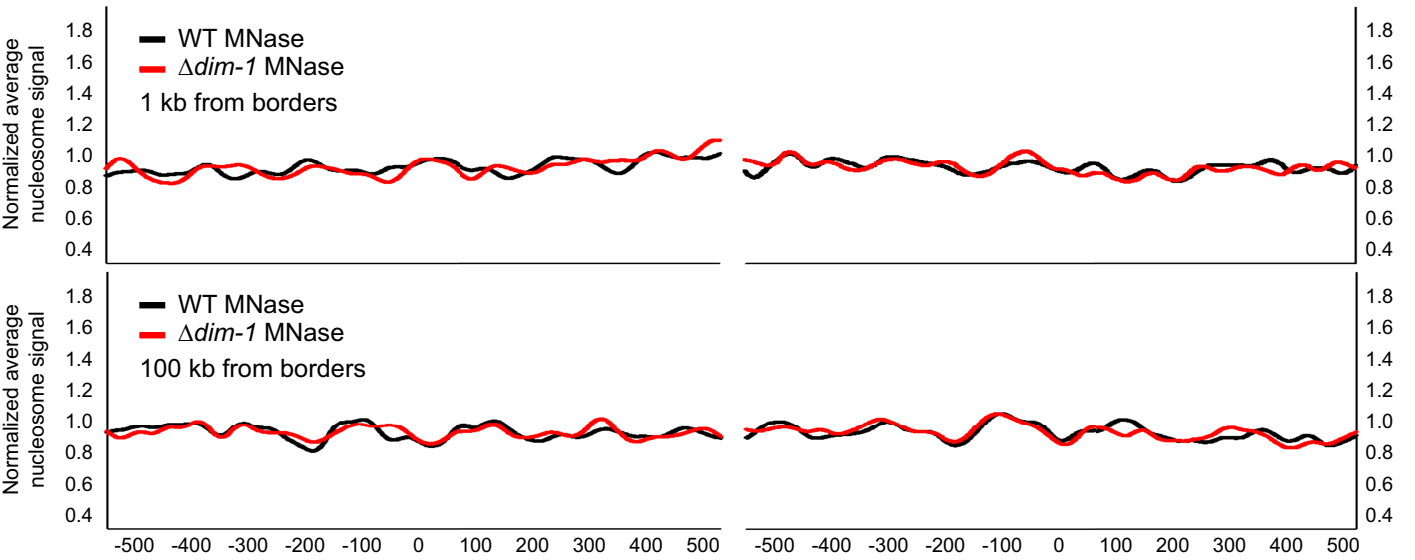

Figure S17. Nucleosomes have minimal periodicity when using random genomic reference points. Average nucleosome enrichment profiles of MNase-seq signal from WT (black) and Δ*dim-1* (red) strains relative to random genomic sites (n = 210) from upstream (left) or downstream (right) of heterochromatic regions. Plots using sites 1,000bp (top) and 100,000bp (bottom) from borders are shown as controls for the analyses in Figure 4.