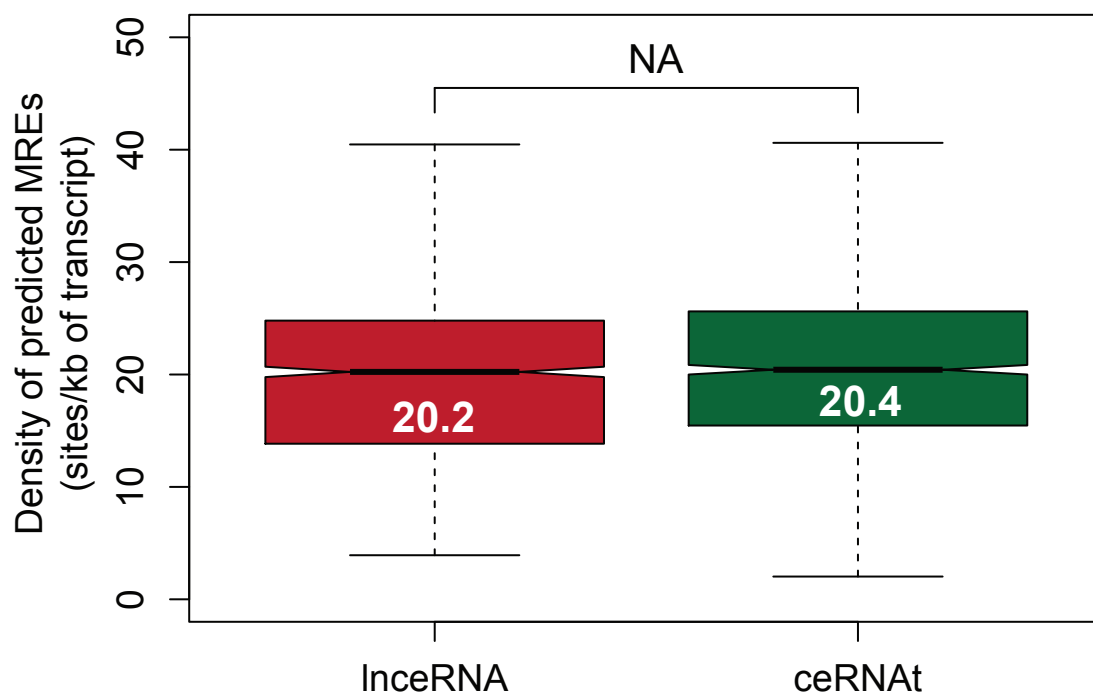




Supplementary Figure S9. Distribution of MREs for lncRNAs and protein-coding genes.

Distribution of the density of predicted MREs for the top 25% most highly expressed miRNAs within lncRNAs (median=20.2 MREs/kb, red) and the 3'UTR of their ceRNA (green, median=20.4 MREs/kb). NA, $p=0.34$.