

Supplemental Figure 9. Reproducibility of counts and ratios across technical replicates. Correlation of DNA tag counts (left/blue), RNA tag counts (middle/green) as well as RNA/DNA ratios (right/red) for pairwise comparisons of the three mutant-integrase (MT) library replicates (A) and the three wild-type-integrase (WT) library replicates (B) along the rows. Both the counts and the ratios are calculated from RNA and DNA counts that are summed across ~60 barcodes representing a given candidate enhancer sequence and represented in a given experiment.

