Supplemental Figure S7. Correlation (Pearson/p-rho and Spearman/s-rho) of DNA tag counts (left column), RNA tag counts (middle column) as well as RNA/DNA ratios (right column) for pairwise comparisons of three wild-type-integrase (WT) library replicates (rows). These are for individual barcodes, *i.e.* before summing across barcodes representing the same candidate enhancer sequence.