



Supplemental Figure 6. The overlap among the genes bound by E2F4 in multiple mouse tissues does not depend on cutoff chosen. Each panel represents the overlap in binding data between the tissues located in the row-column combination, as labeled at p-values of (A) 1×10^{-3} , (B) 1×10^{-4} , and (C) 1×10^{-5} . The total number of genes bound in each tissue, given the p-value cutoff, is shown as the white background diagonal that represents the same tissue row-column (e.g. acinar versus acinar, et cetera).