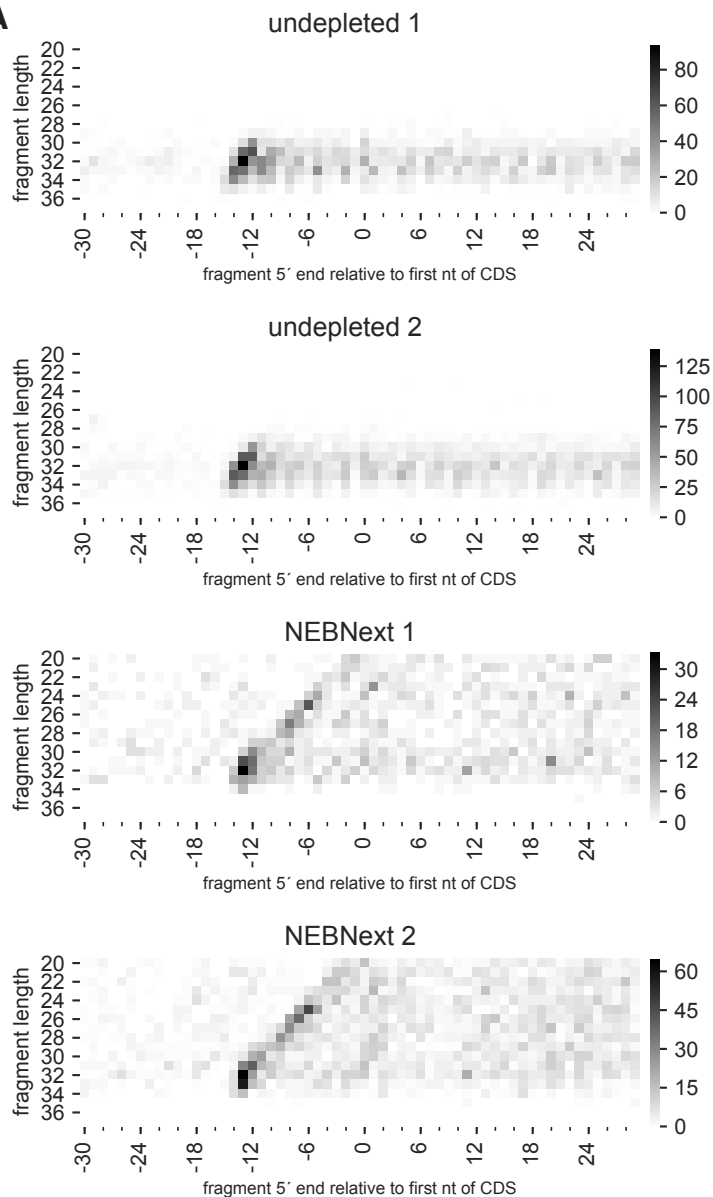
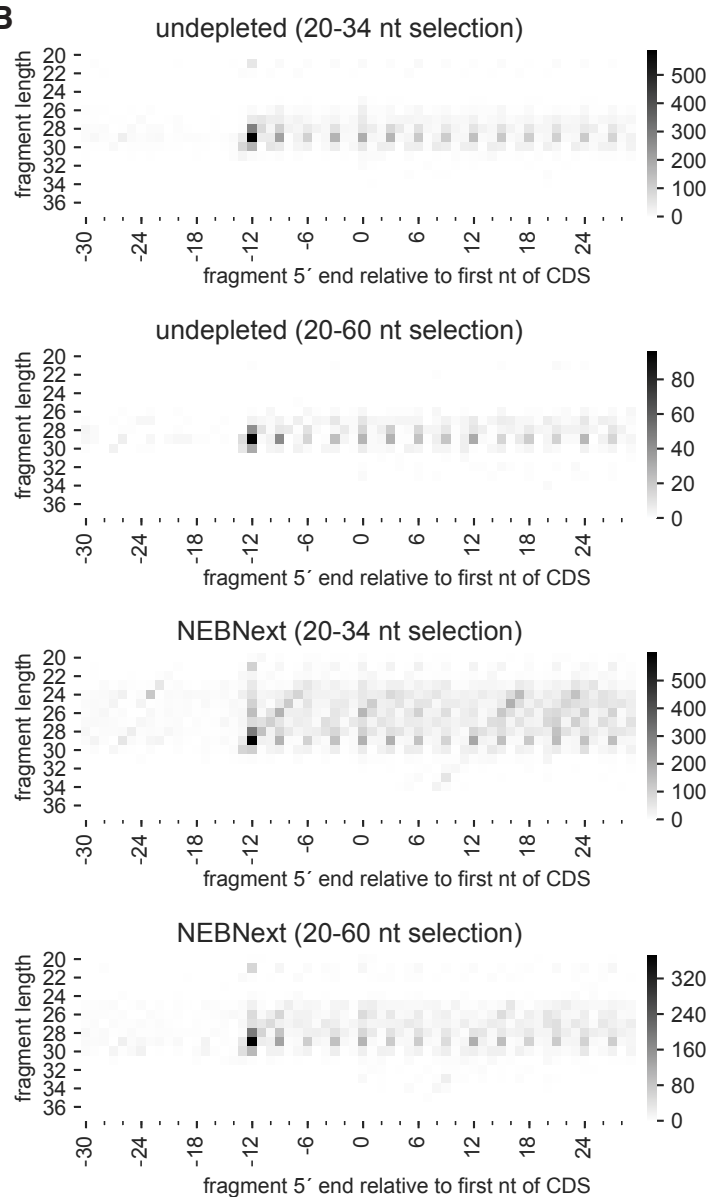
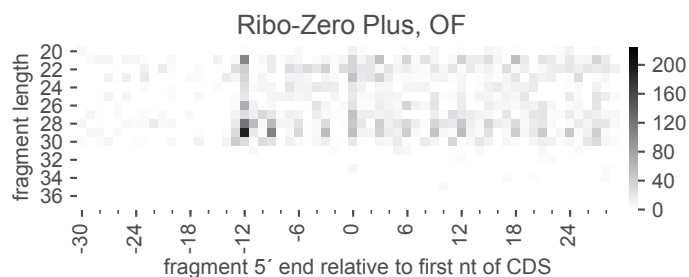
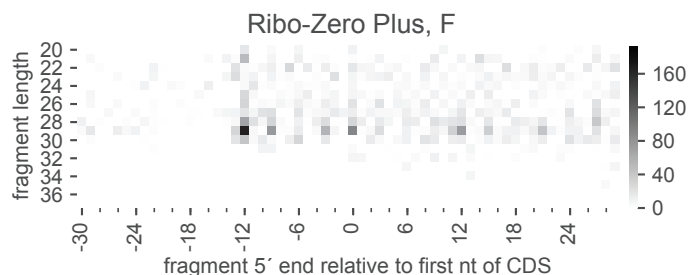
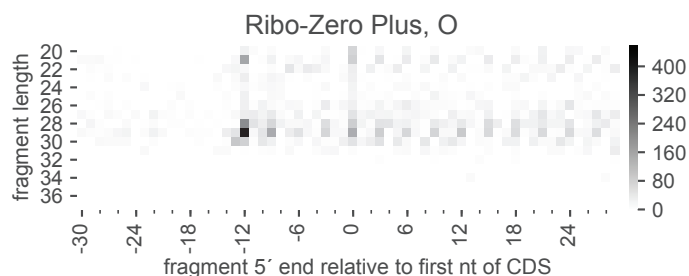
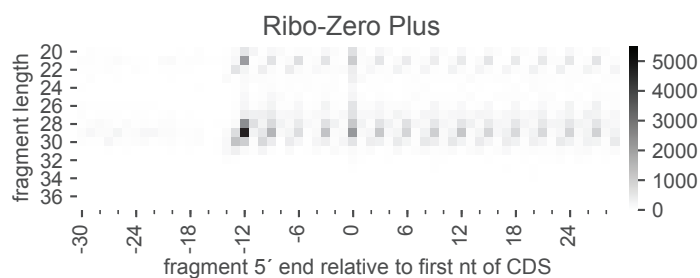
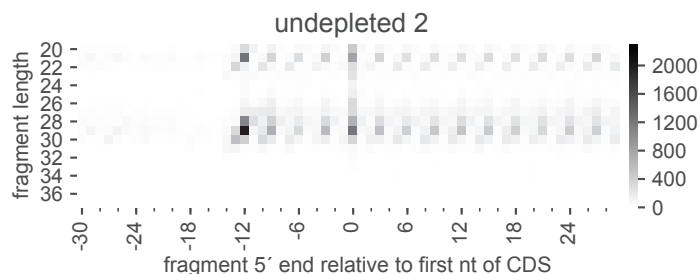
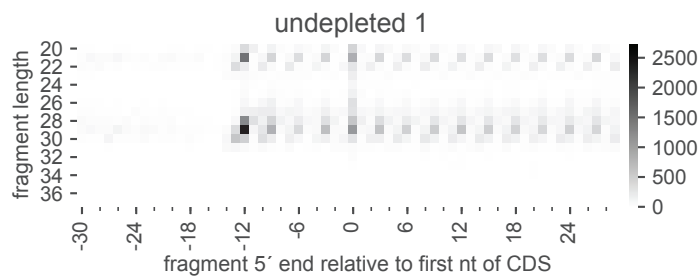


A**B**

Supplementary Figure 1: Fragment 5' ends mapping to CDS starts by read length for NEB-Next depletion.

Heatmaps of number of ribosome footprint 5' ends mapping relative to the first nucleotide of coding regions, subdivided by read length, for NEBNext depleted libraries from A) Figure 1A and B) Figure 1C.



Supplementary Figure 2: Fragment 5' ends mapping to CDS starts by read length for Ribo-Zero plus depletion.

Heatmaps of number of ribosome footprint 5' ends mapping relative to the first nucleotide of coding regions, subdivided by read length, for Ribo-Zero Plus depleted libraries from Figure 1E.