SUPPLEMENTAL FIGURE S2. Distribution of 7-mer enrichment values in bind-n-seq experiments involving PPR10 variants. Enrichment values were calculated for each 7-mer at each position as the frequency of that 7-mer in the bound fraction divided by its frequency at the same position in the input library. The subset of 7-mers that were enriched more than 5 or 10 standard deviations above the mean are marked.