



Figure S4: PRO-cap signal in *S. cerevisiae* is consistent with characterized TATA box and TFIIB positions. **A.** Median PRO-cap signal for each base is plotted relative to previously characterized TATA box, or TATA-like motif, coordinates. **B.** Median PRO-cap data at each base relative to TFIIB position for TATA-containing and TATA-like gene groups. Gene groups, as well as motif and TFIIB coordinates were identified previously (Rhee and Pugh 2012). The y-axis for all plots shows the median read counts for each base. The 12.5% and 87.5% quantiles are shown in lightly shaded regions.