Figure S2: Comparison of PRO-cap-observed transcription start sites with previous annotations of *S. pombe* TSSs (Eser et al. 2016). A. Box plots showing the distribution of distances to reannotated TSSs from previous positions (positive values are downstream, negative values are upstream of previous annotations). B. Composite PRO-cap signal intensity (background subtracted) centered on each set of TSS annotations for the same genes (N = 2646).