



**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).