



Figure S2 Gene-specific examples of ORF nucleosome shifts. UCSC Genome Browser screenshots showing the distribution of scaled MNase-Seq read counts for two biological replicates (alpha and beta) of wild-type cells grown in YPD (Glu WT), cells not expressing Yta7 (Glu *yta7*), wild-type cells grown in YPGal (Gal WT) and cells over-expressing Yta7 (Gal *YTA7oe*). *yta7* cells exhibited a 5' shift in nucleosome position, whereas *YTA7oe* cells exhibited a 3' shift in nucleosome position relative to WT, consistent with metagenome analysis.