



**Supplemental Figure S2: (A-C)** Analysis of the fraction of CPDs remaining following 2hr repair adjacent to the transcription start site (TSS) from CPD-seq data for WT, *snf5* $\Delta$ , and *snf6* $\Delta$  cells. Average NCP coverage indicates the read coverage for nucleosome dyads identified in a previous study (Weiner et al. 2015).