



Figure S3 Repair activity is significantly different across chromatin types genome-wide. (A,B)

Distribution of repair activity (30min) for bins of each chromatin type in the Chromosome 2R subset from figure 2C (A) and genome-wide (B). Data is normalized to cellular damage levels. The letters above each plot categorize statistically different distributions, as determined by a one-way ANOVA and Tukey's multiple comparisons test ($p < 0.01$). Outliers beyond the whiskers of each plot represent the bottom 2.5% and top 2.5% of that distribution.