



Supplementary Figure 3. Fragment Enrichment Over Gene Body. Plotting peak enrichment over the gene body for all time points (0, 18, 48, and 120 hours post-RA) and biological replicates (rep1 or rep2) which showed the most accessibility occurs around the transcription start sites. Accessibility is noted here in terms of fragment density of transposition events along genes within the genome. Peaks at 0bp correspond with the either the transcription start sites (TSS) or transcription end site (TES) of any given gene in the genome.