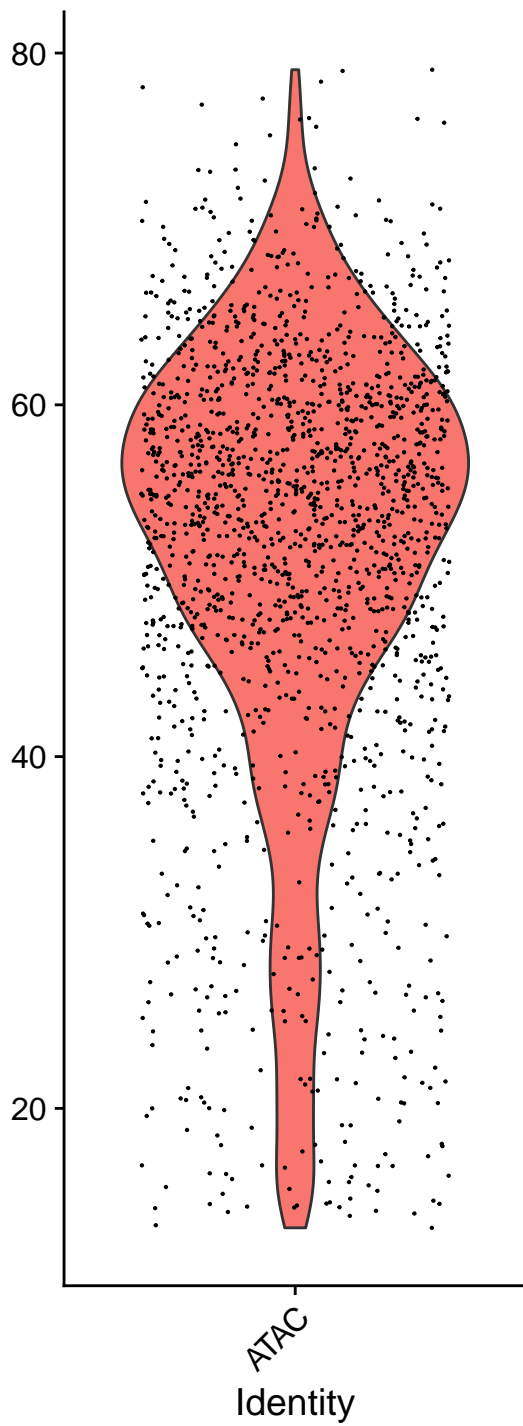
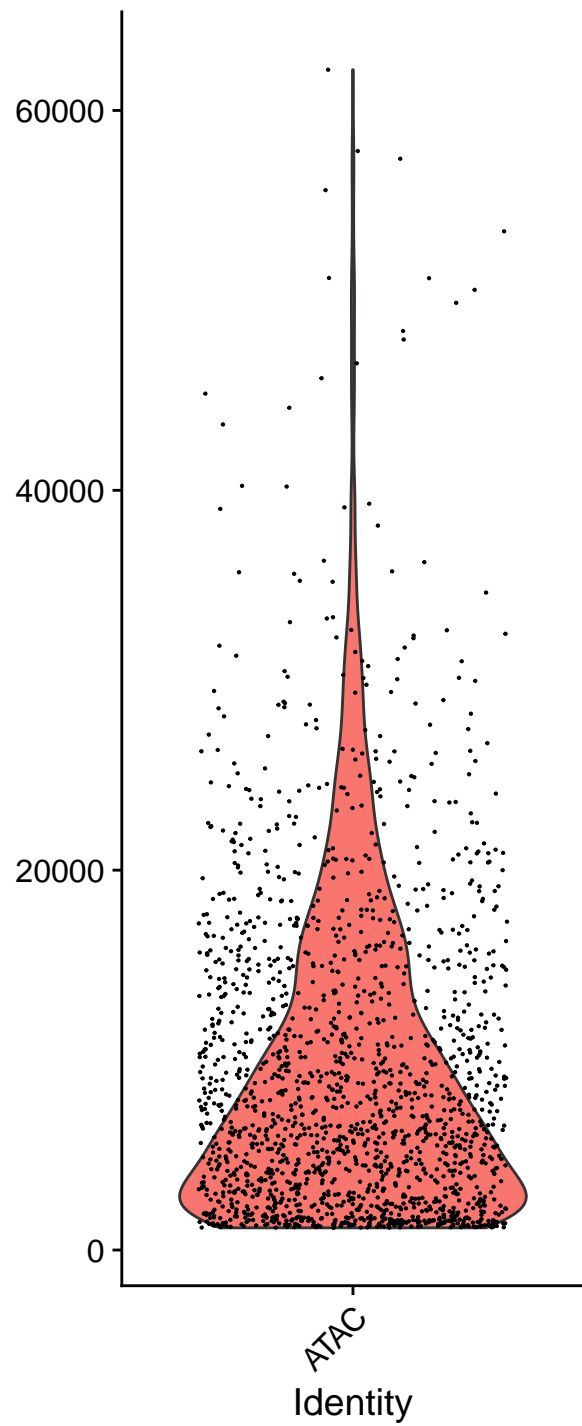


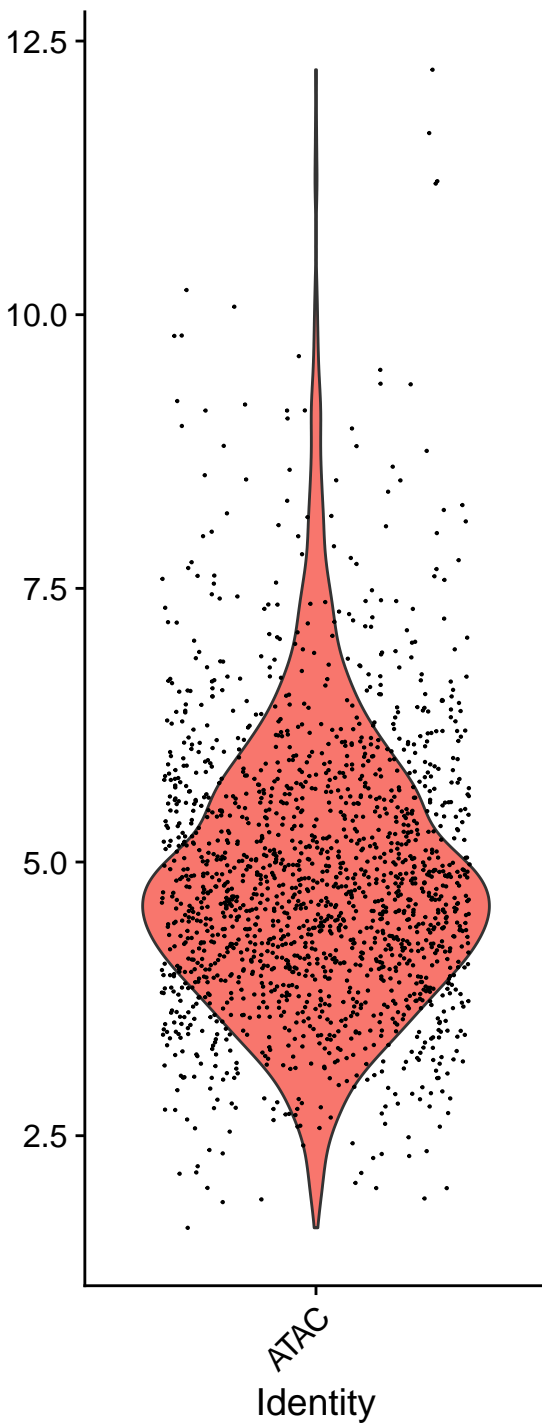
pct_reads_in_peaks



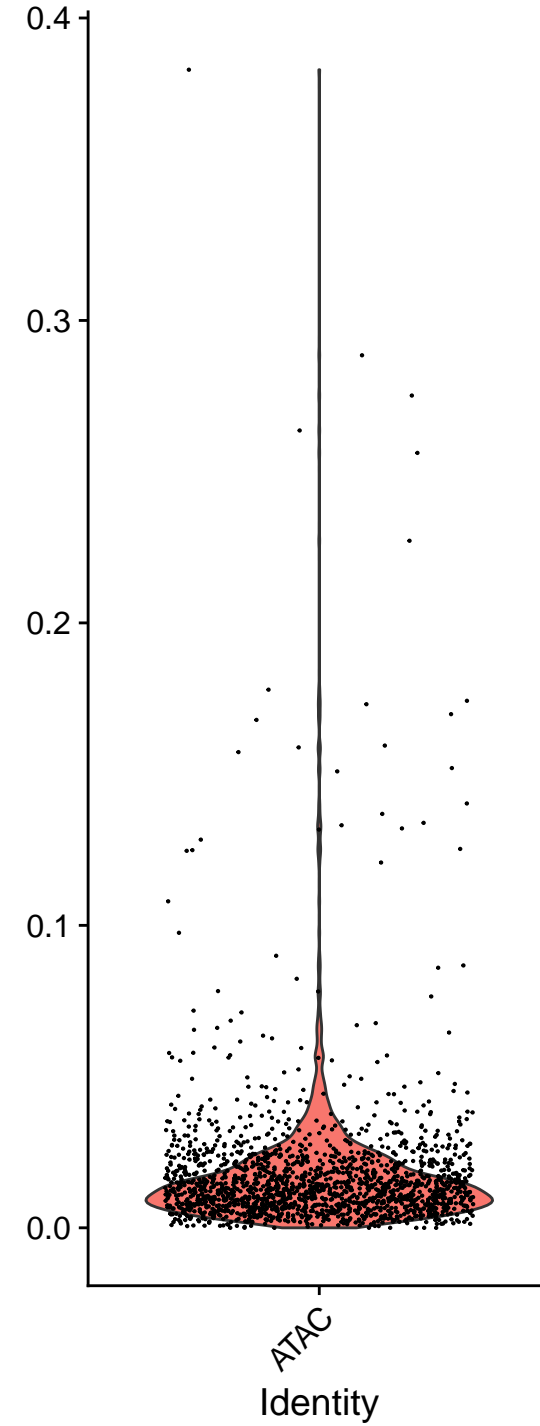
peak_region_fragments



TSS.enrichment



blacklist_ratio



nucleosome_signal

