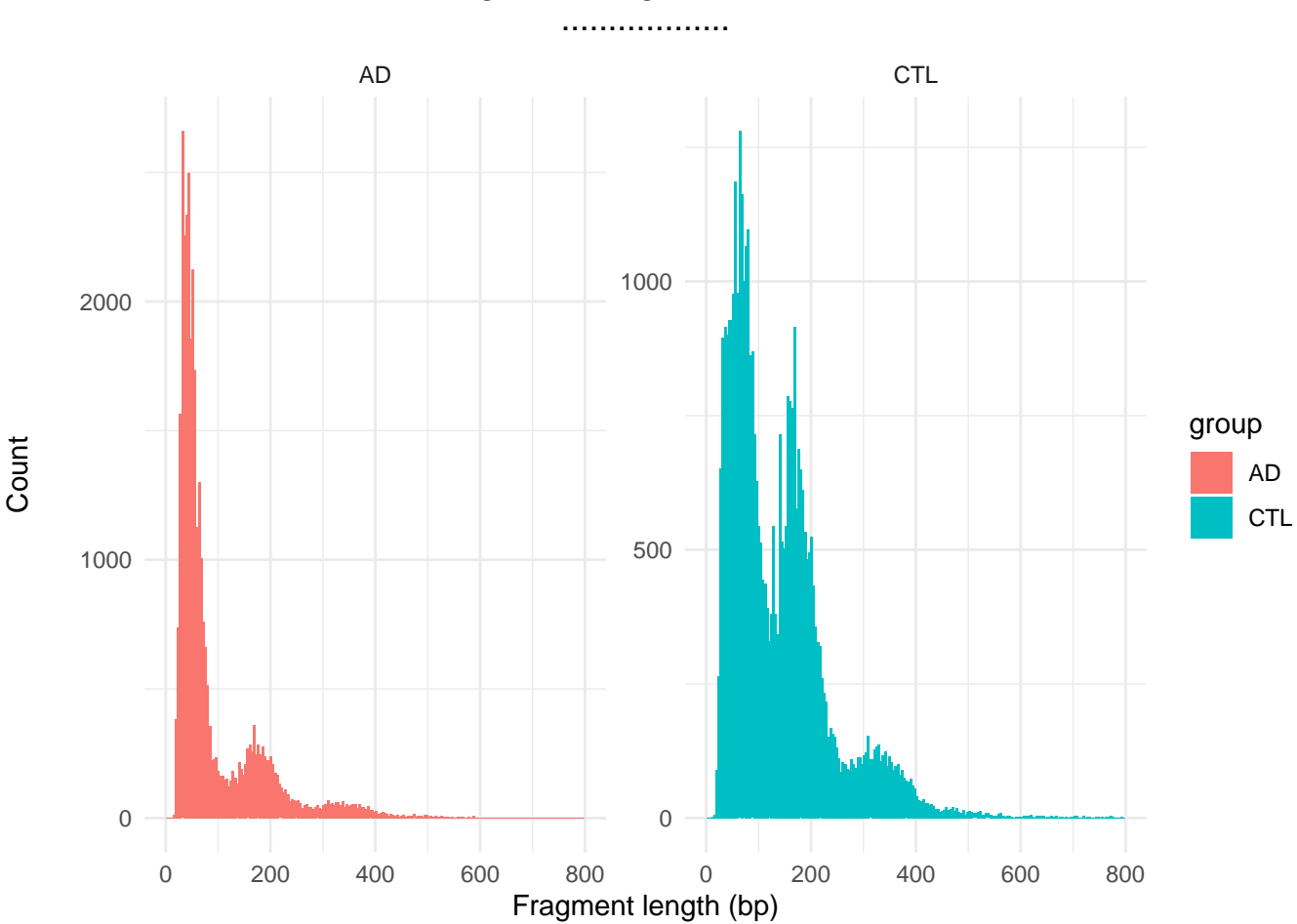
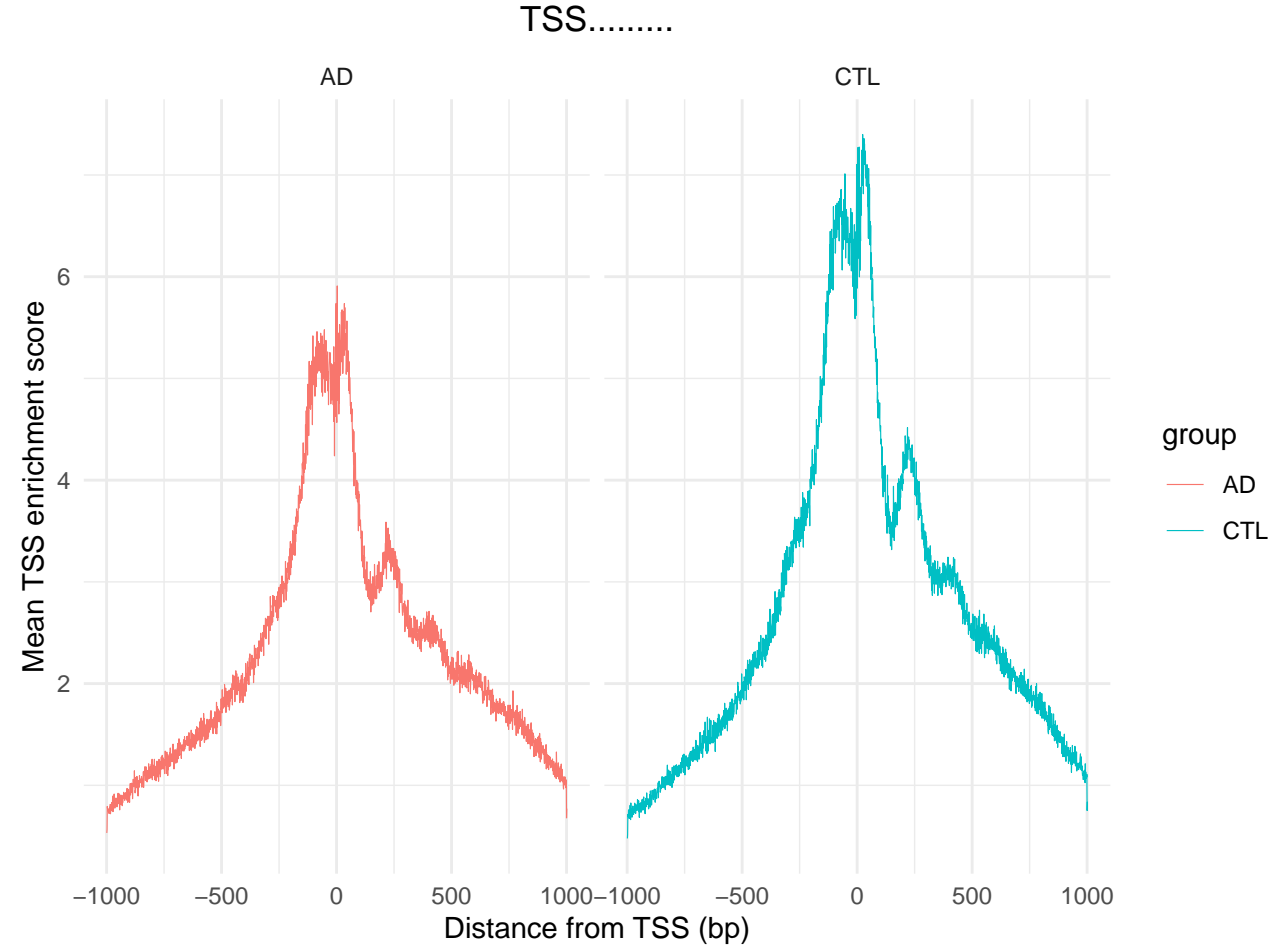


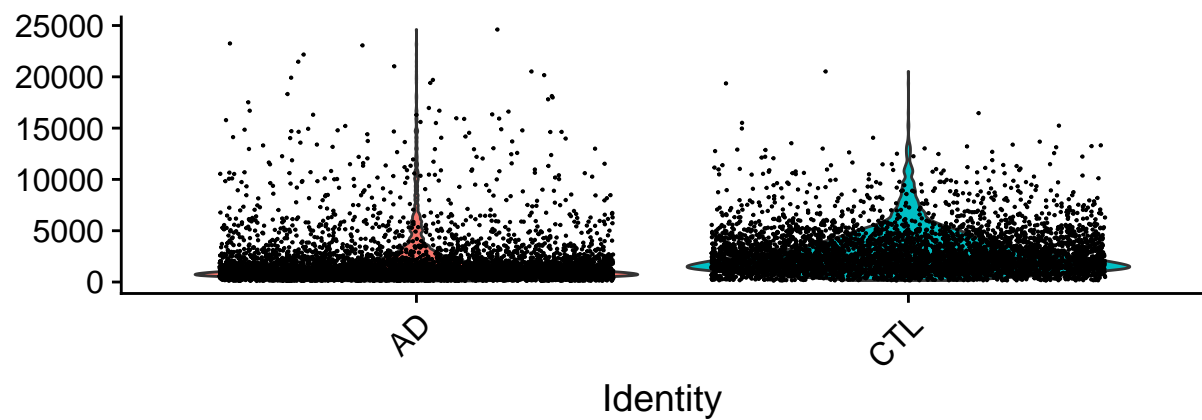
Fragment Length Distribution



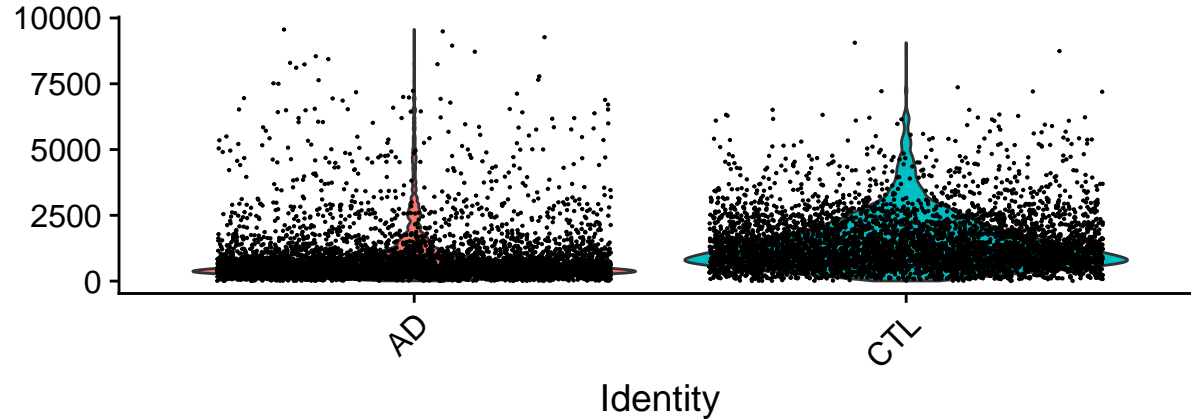
TSS Enrichment Profile



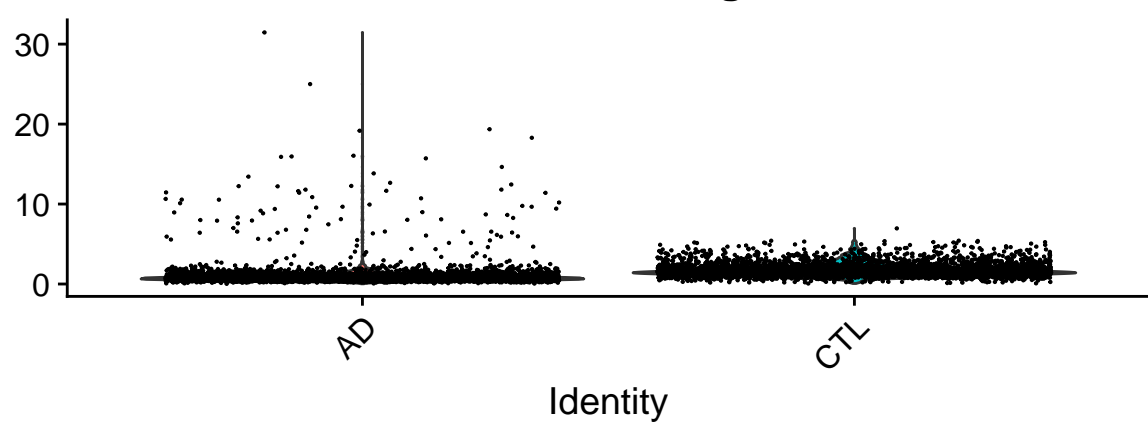
nCount_peaks



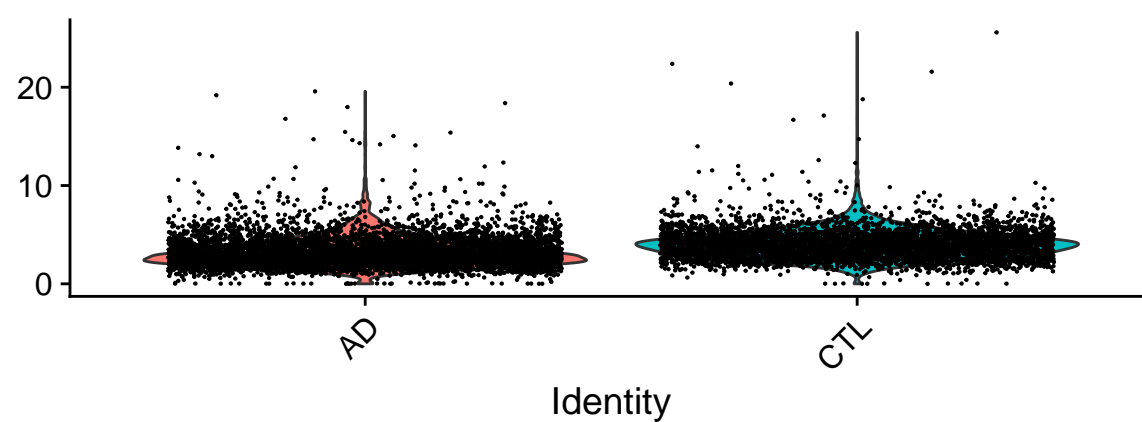
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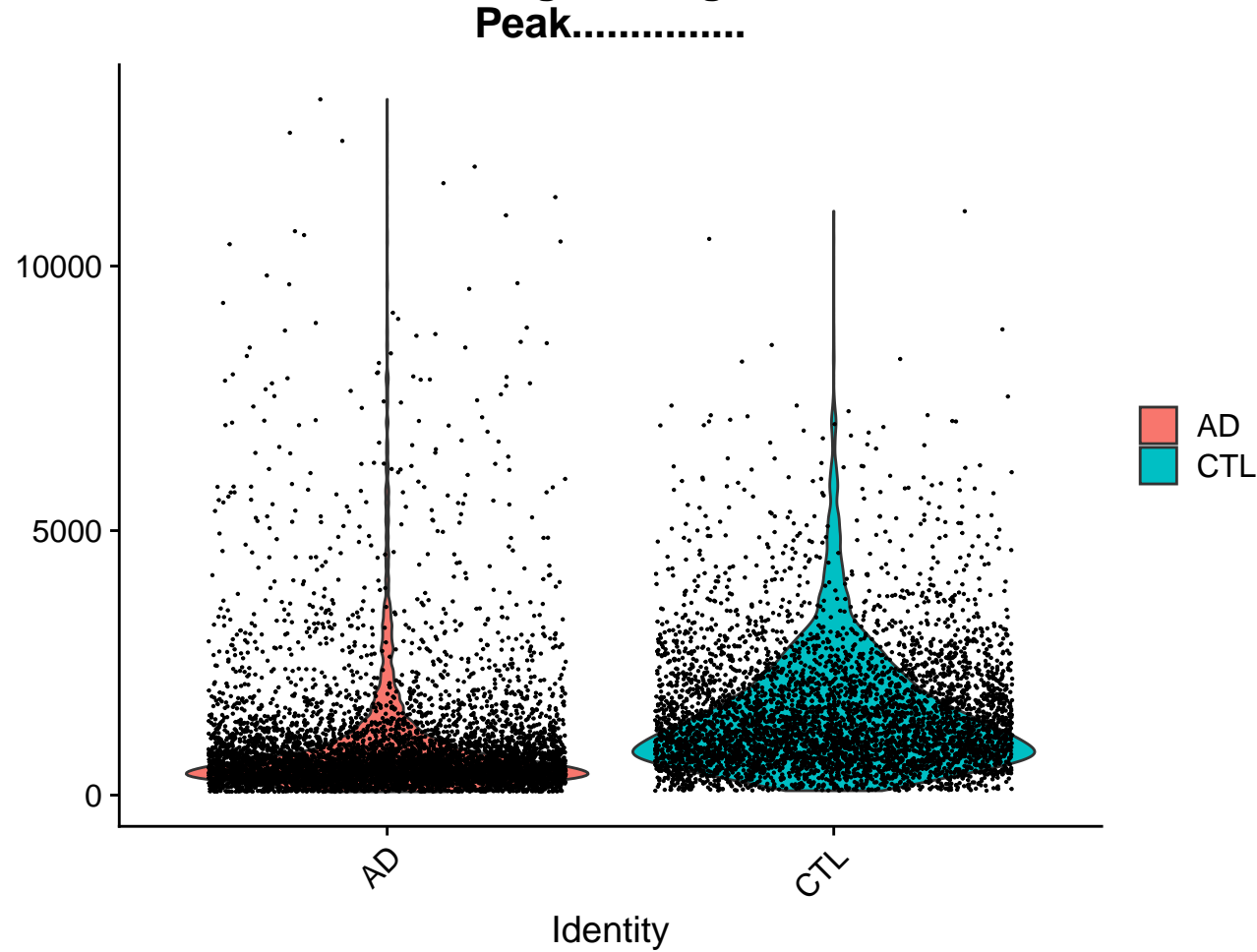
nucleosome_signal



TSS.enrichment



Peak Region Fragments



Blacklist Ratio

