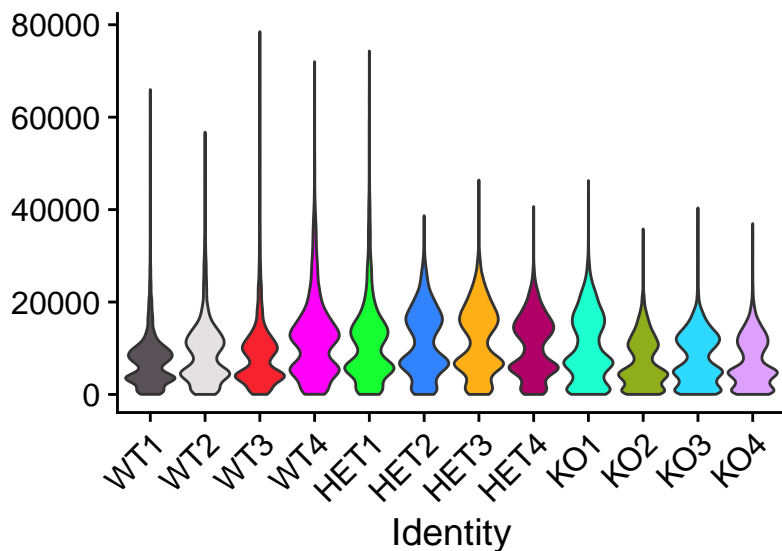
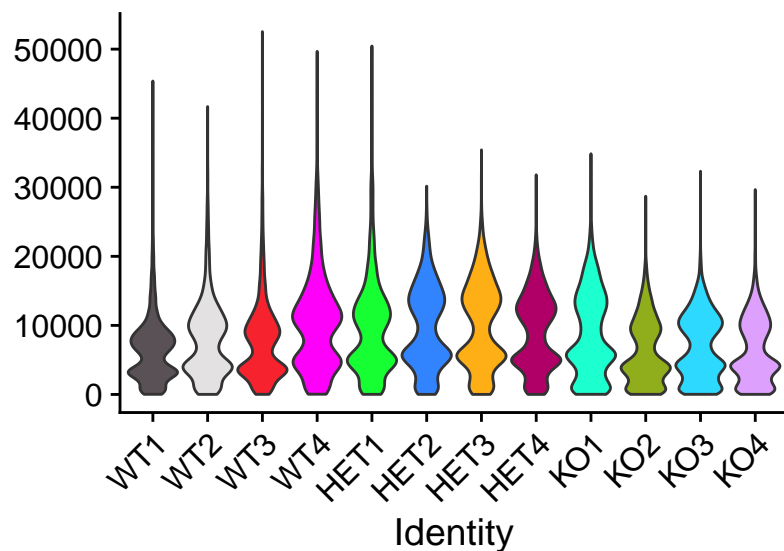


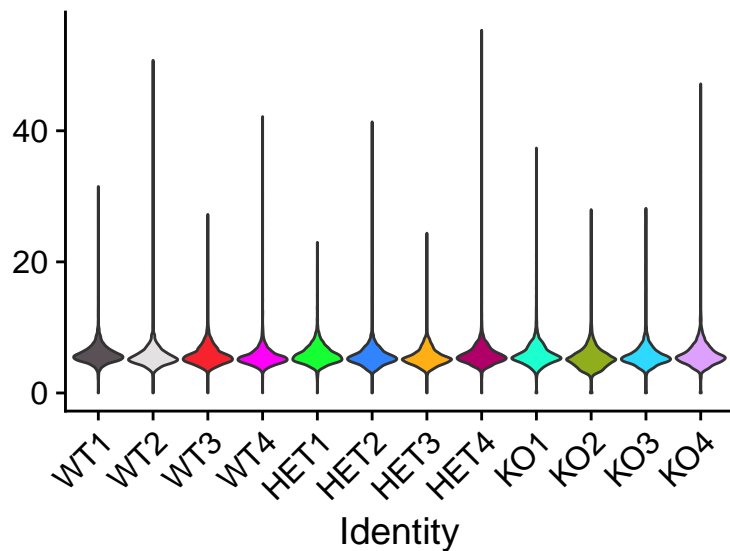
**nCount\_ATAC**



**nFeature\_ATAC**



**TSS.enrichment**



**nucleosome\_signal**

