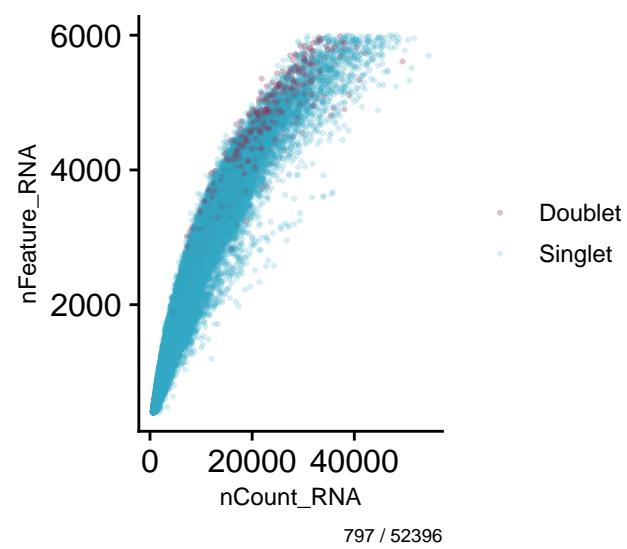
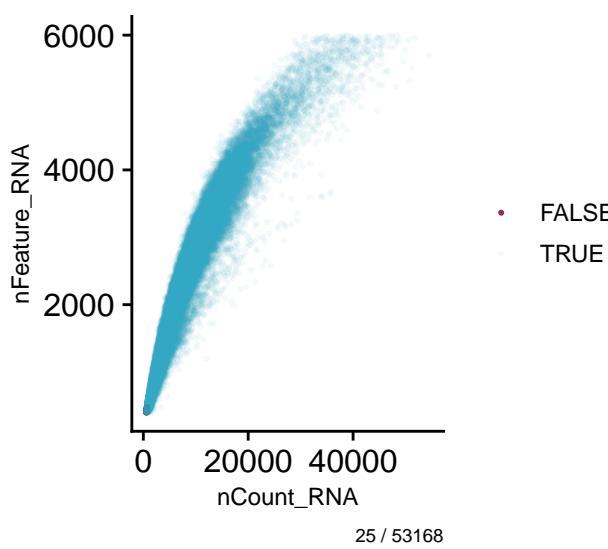


## QC including scran and two doublet methods intersection

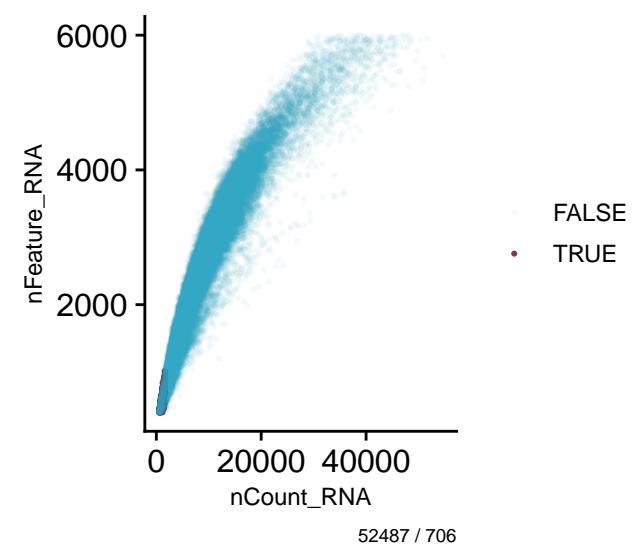
**DOUBL\_INTERSECT**



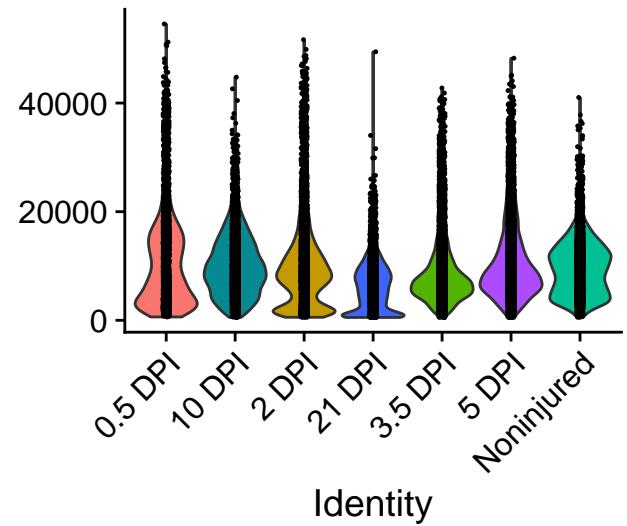
**is\_cell**



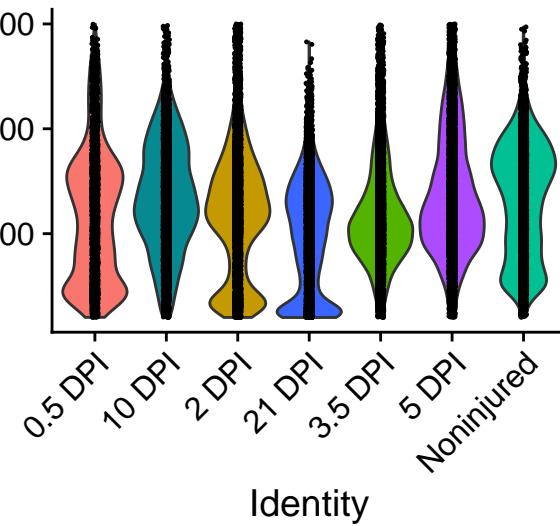
**is\_inf\_outlier**



**nCount\_RNA**



**nFeature\_RNA**



**percent.mt**

