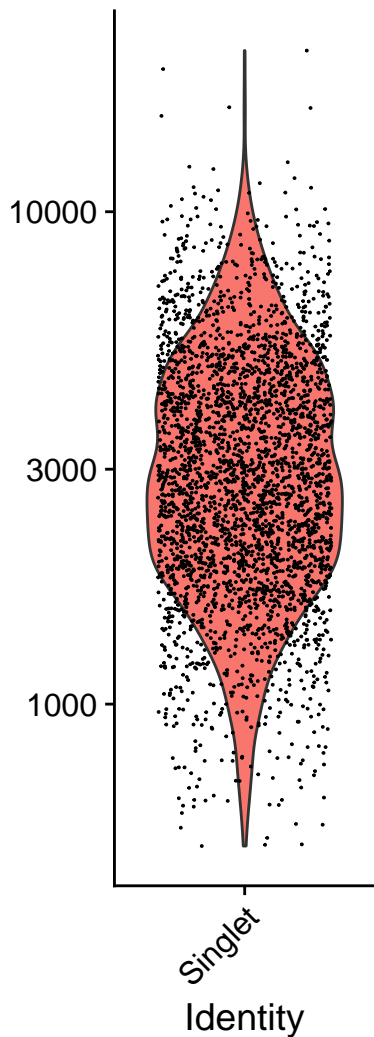
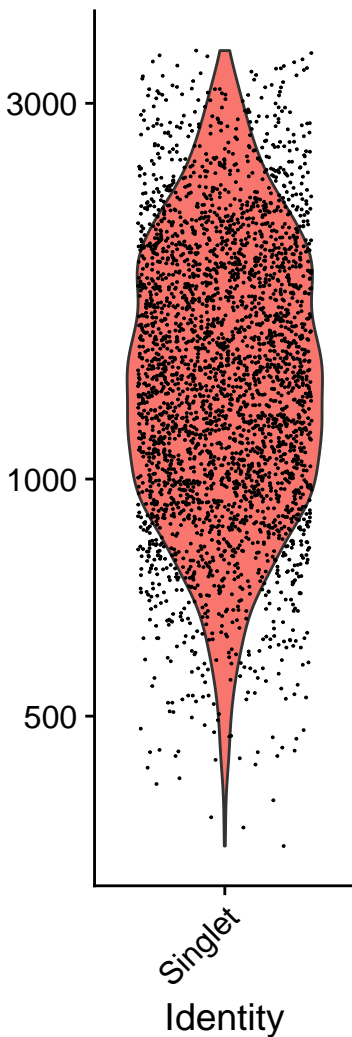


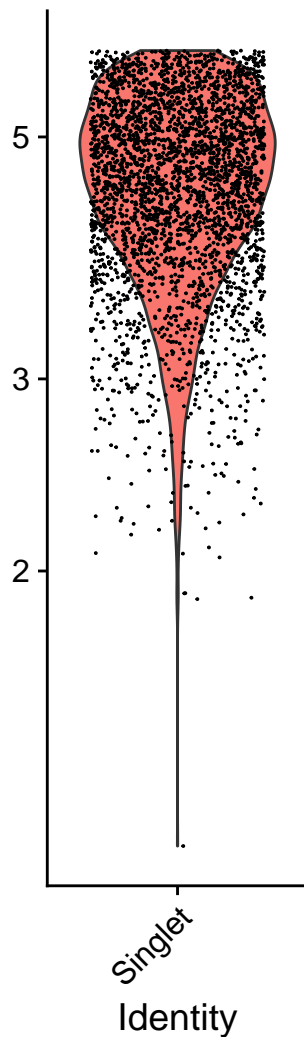
nCount_RNA



nFeature_RNA



percent.mt



QC plot

Number of detected genes in function of number of UMI

