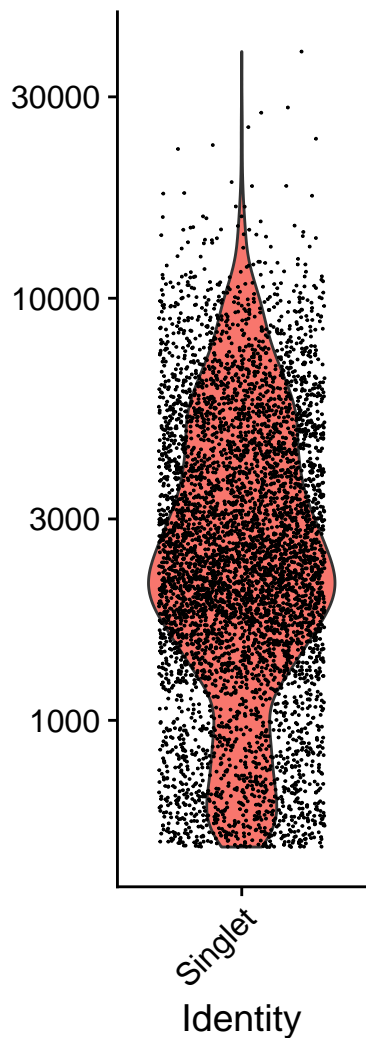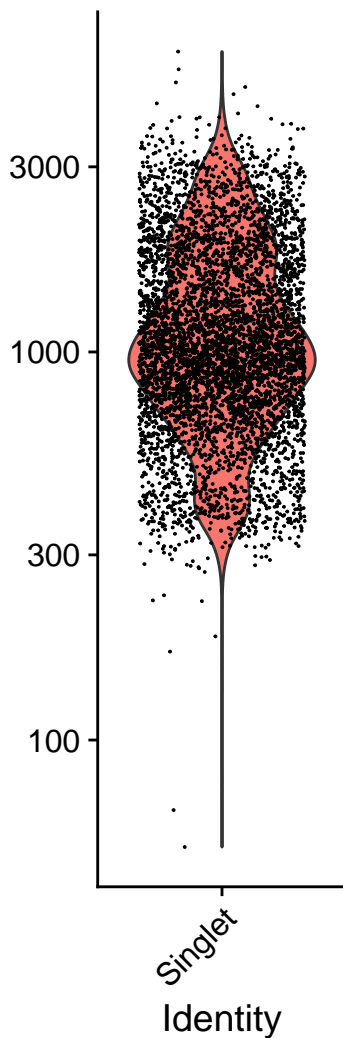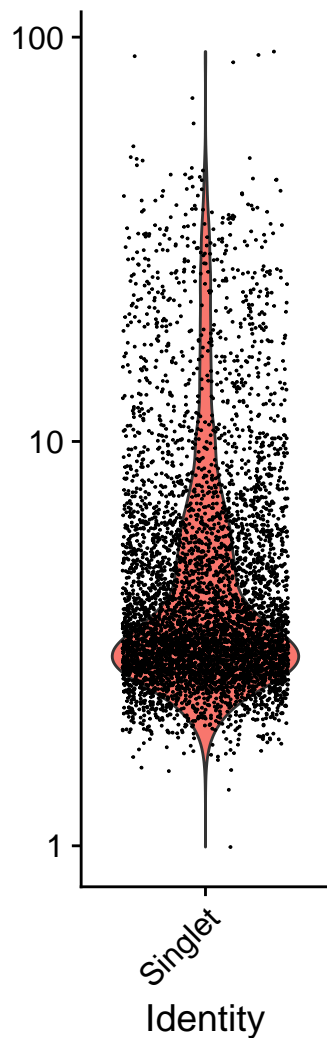| nCount_RNA | nFeature_RNA | percent.mt |

QC plot

Number of detected genes in function of number of UMI