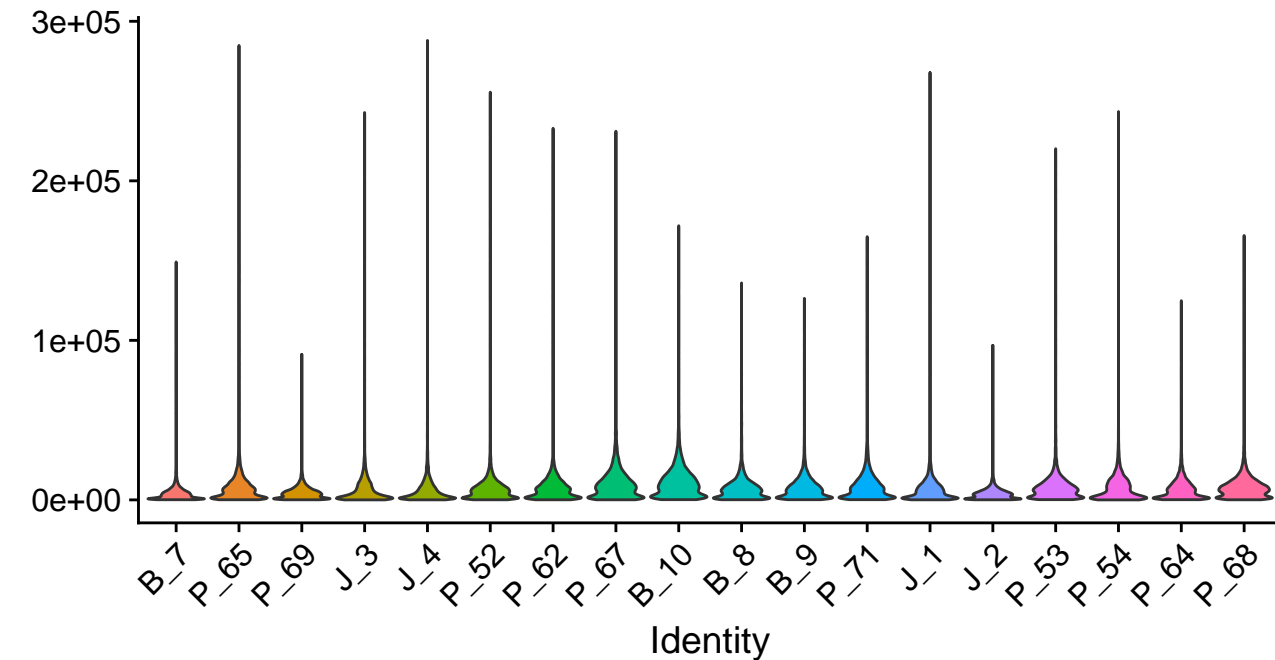
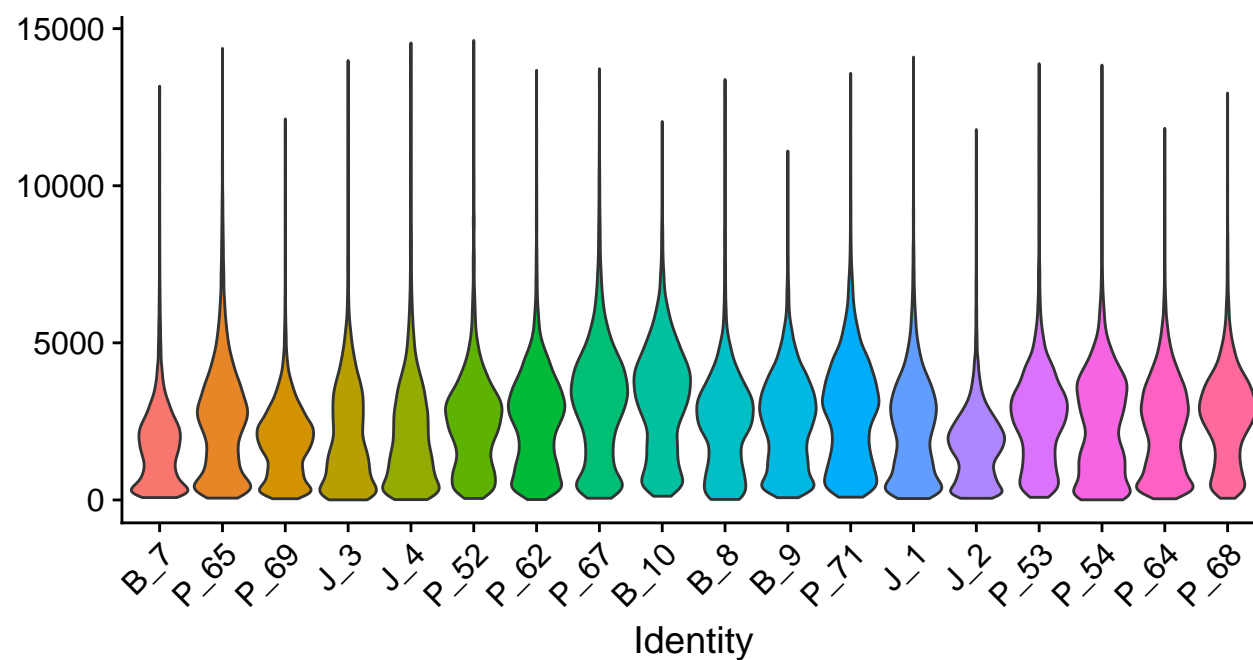


nCount_RNA**nFeature_RNA****cell.complexity**