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Diagnostic plots for introduce referenced seurat -> number of genes detected per cell - violin plot: violinNGene  
width: 300 number of UMIs per cell - violin plot: violinNUMI  
height: 300 percentage of mitochondrial genes per cell - violin plot: violinMito  
nGene and nUMI - scatter plot: scatterNUMINGene  
nUMI and percent.mito - scatter plot: scatterNUMIMito
```