

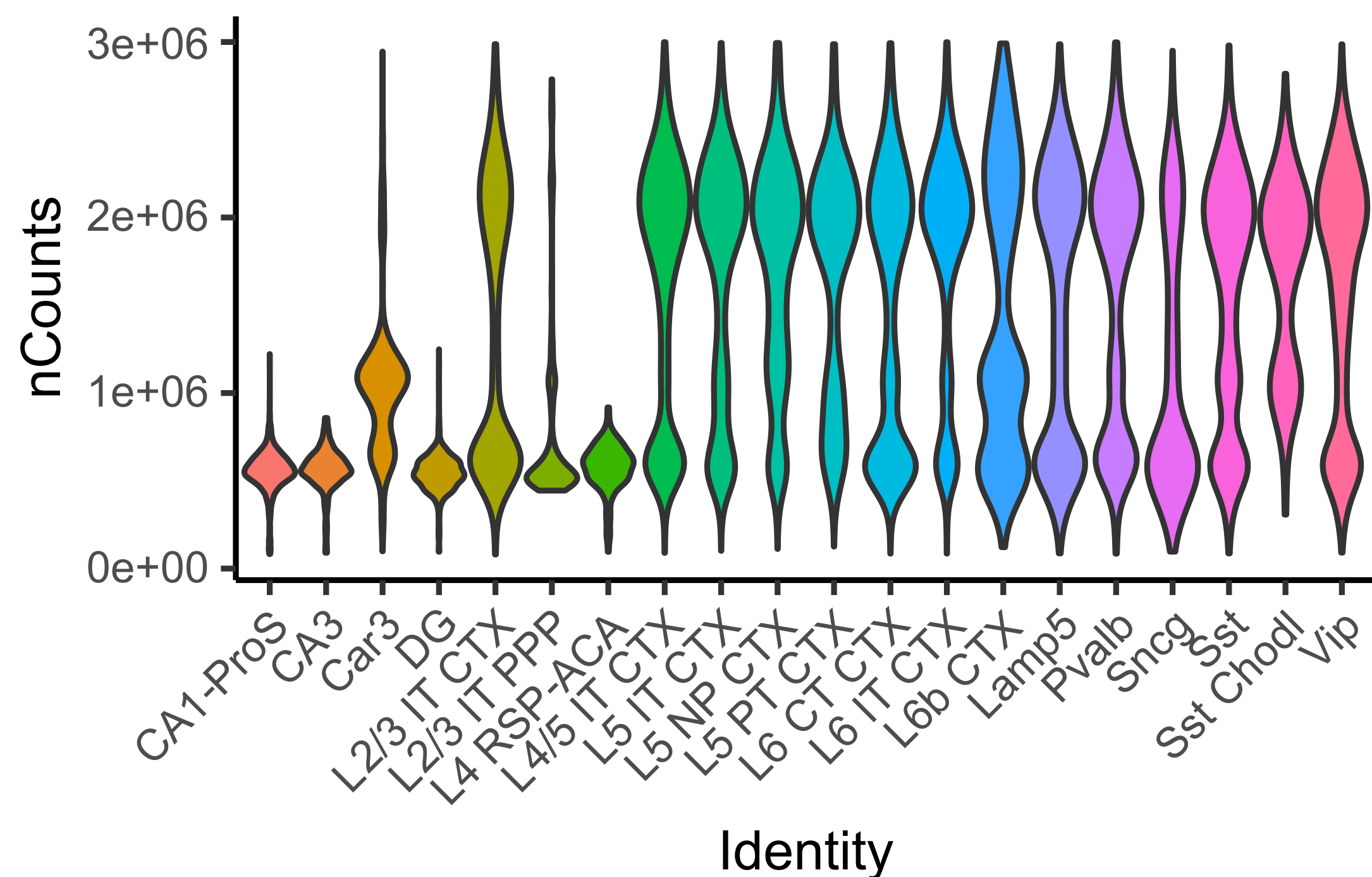
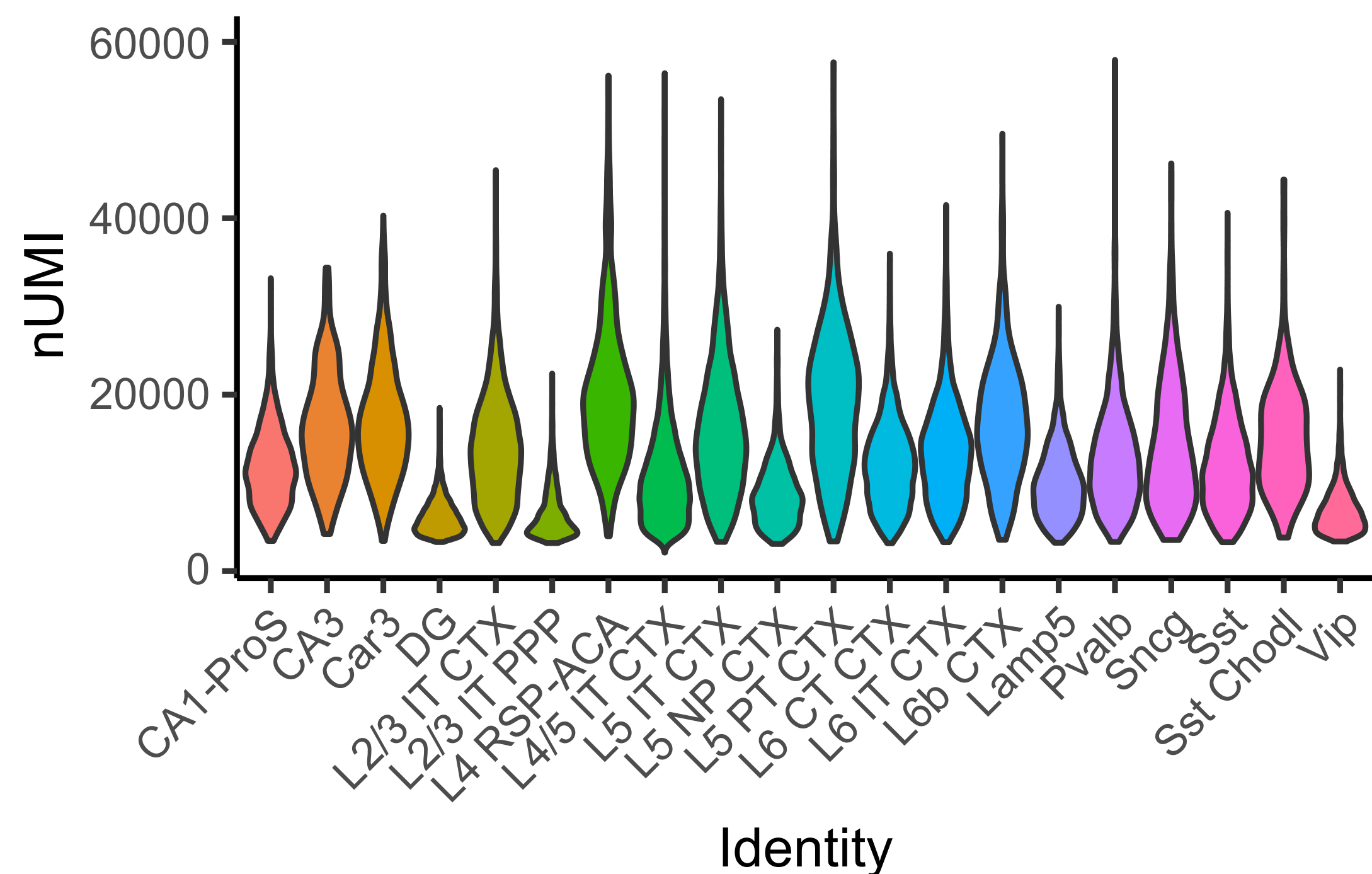
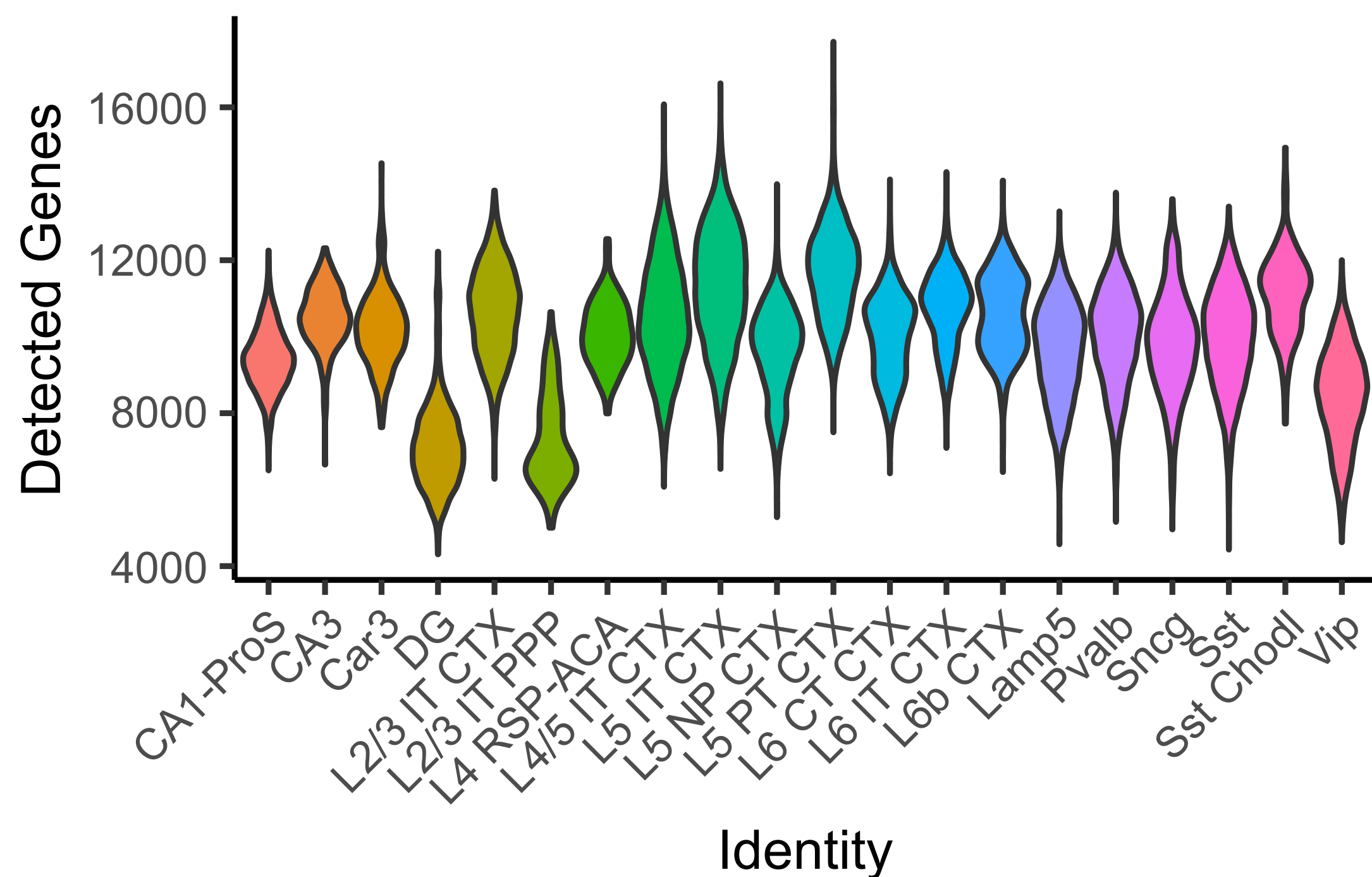
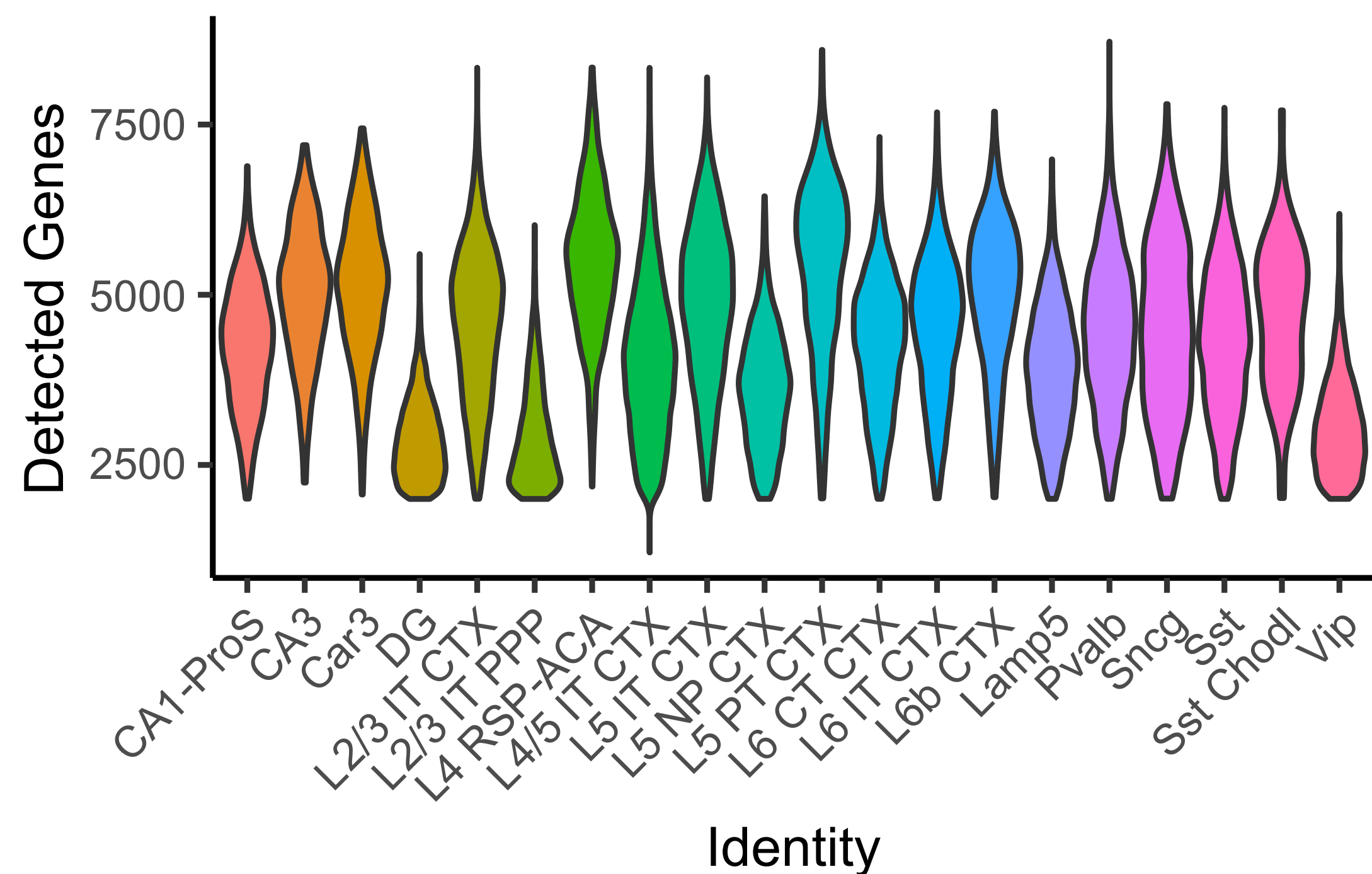
A**Smart-seq Dataset****B****10x Genomics Dataset****C****D**

Figure S1. Quality control metrics for the Smart-seq2 and 10x Genomics datasets. Violin plots showing the distribution of counts/UMIs and detected genes per cell across neuronal subclasses. (A) Number of aligned counts (reads) per cell in the Smart-seq2 dataset. (B) Number of Unique Molecular Identifiers (UMIs) per cell in the 10x Genomics dataset. (C) Number of detected genes per cell in the Smart-seq2 dataset. (D) Number of detected genes per cell in the 10x Genomics dataset.