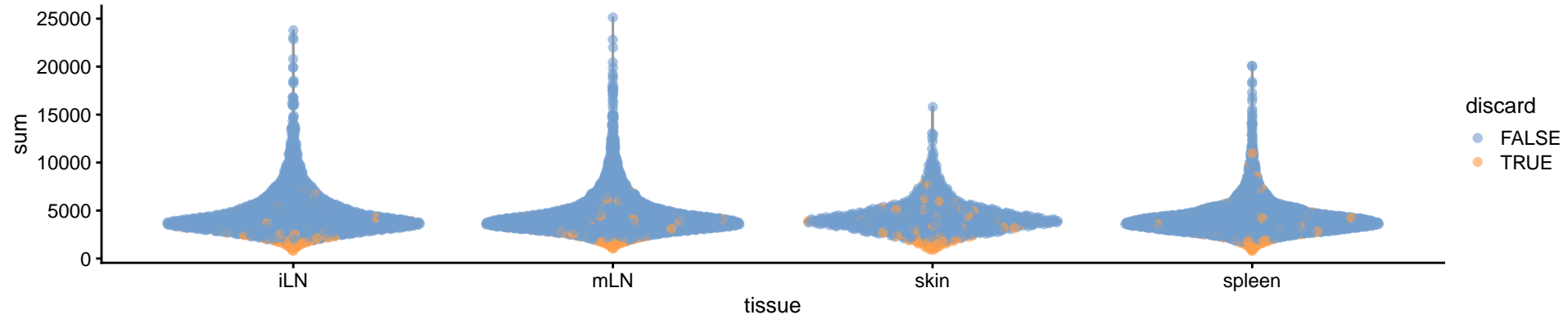
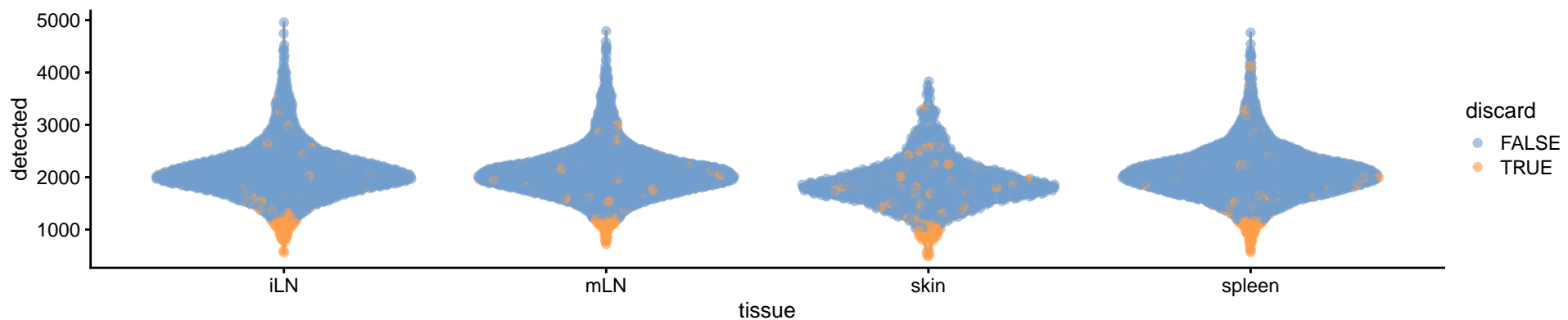


**Sum of all counts (library size)**



**Number of detected genes**



**Percent mitochondrial genes**

