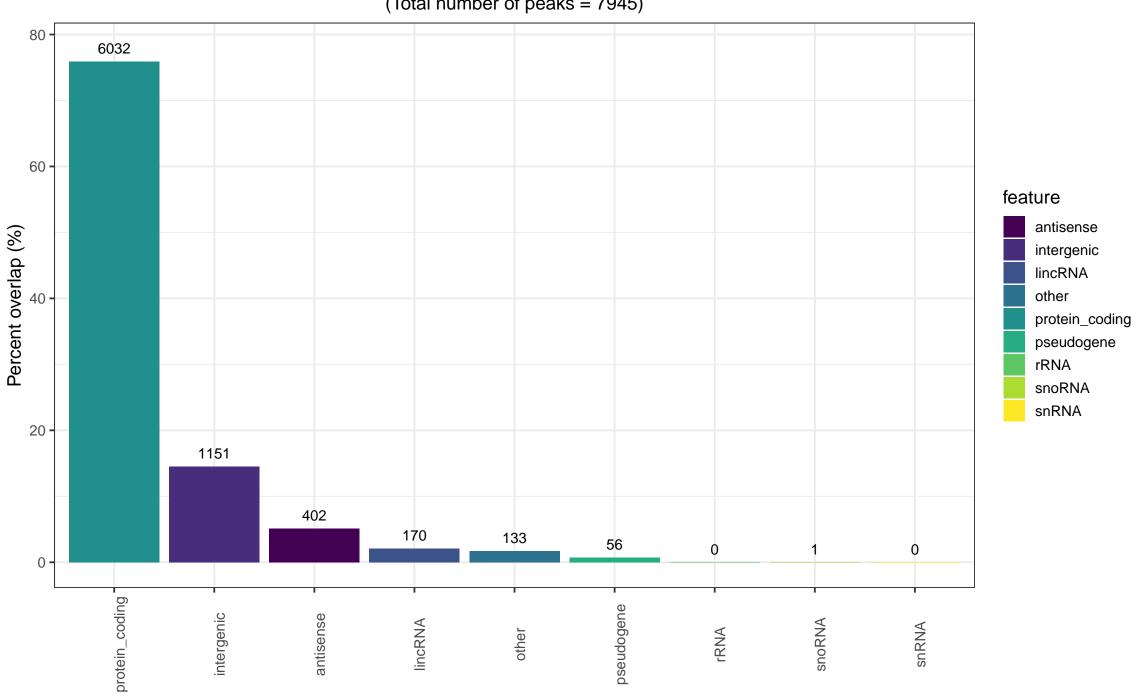
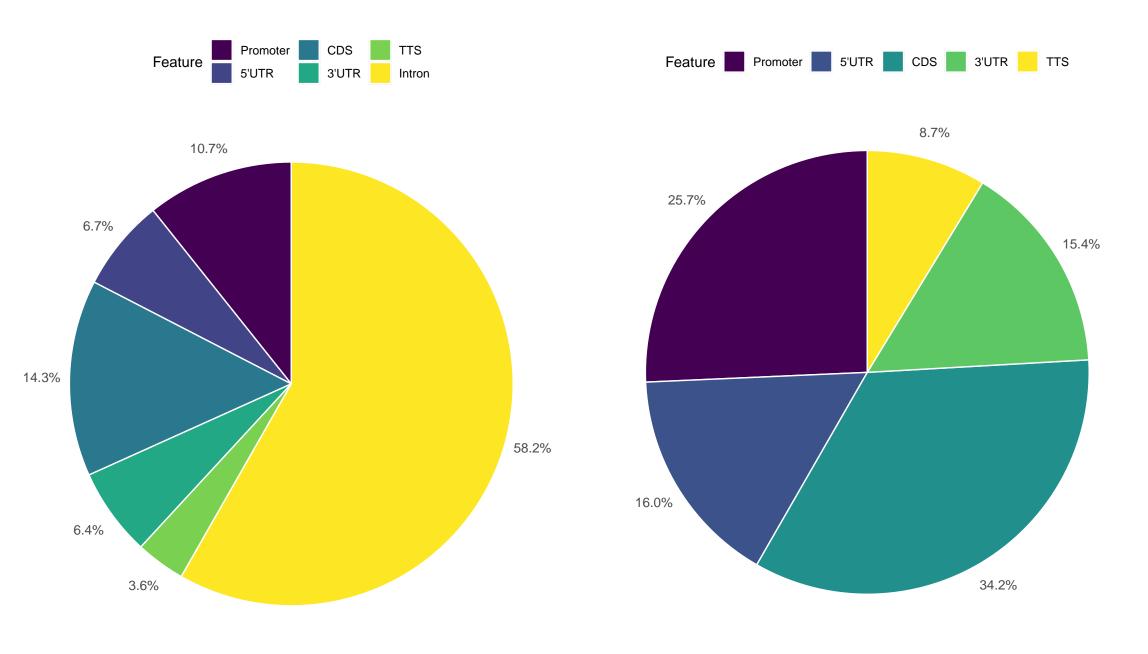
## Annotation of peaks to all type of genes

(Total number of peaks = 7945)



## **Absolute counts**



## Length-normalized counts

