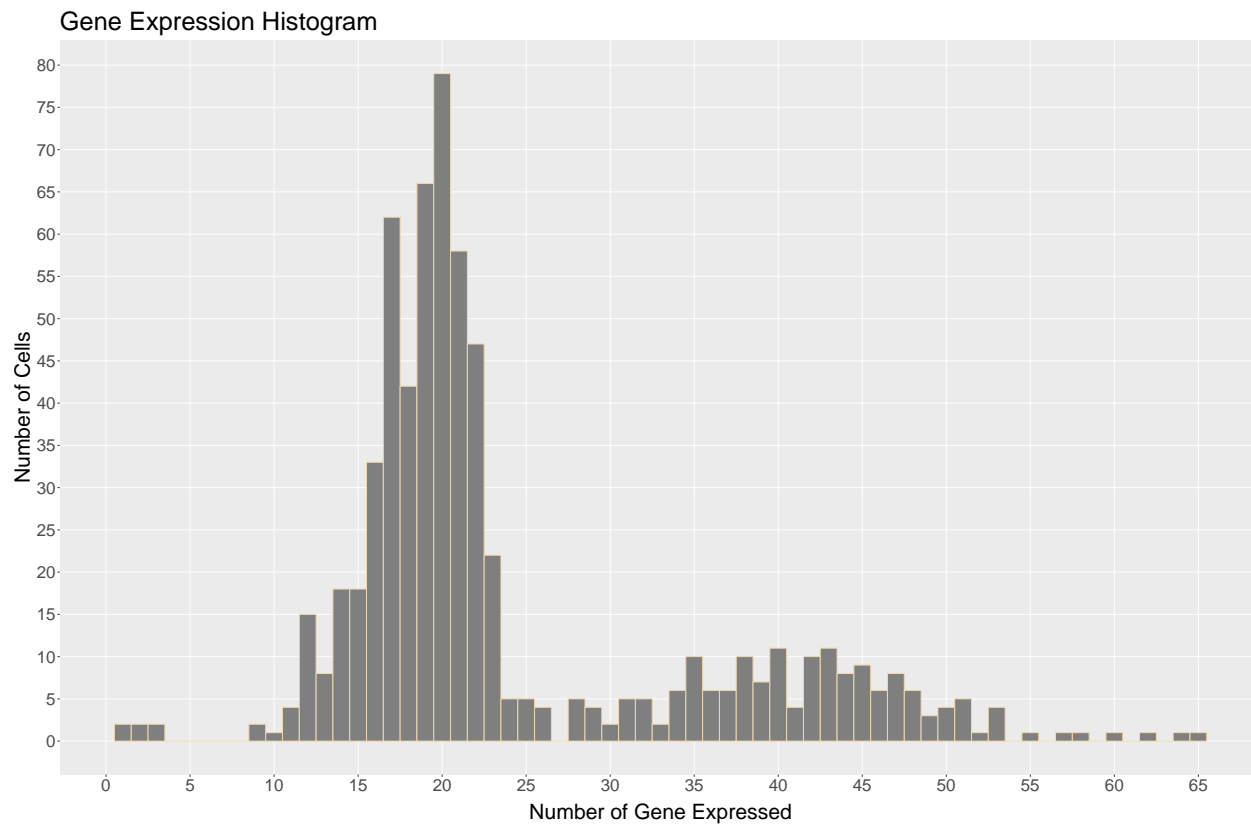


# T1D\_specific\_analysis

No expression detected in 5/1291 cells

```
ctNorm[is.na(ctNorm)] <- 0
```

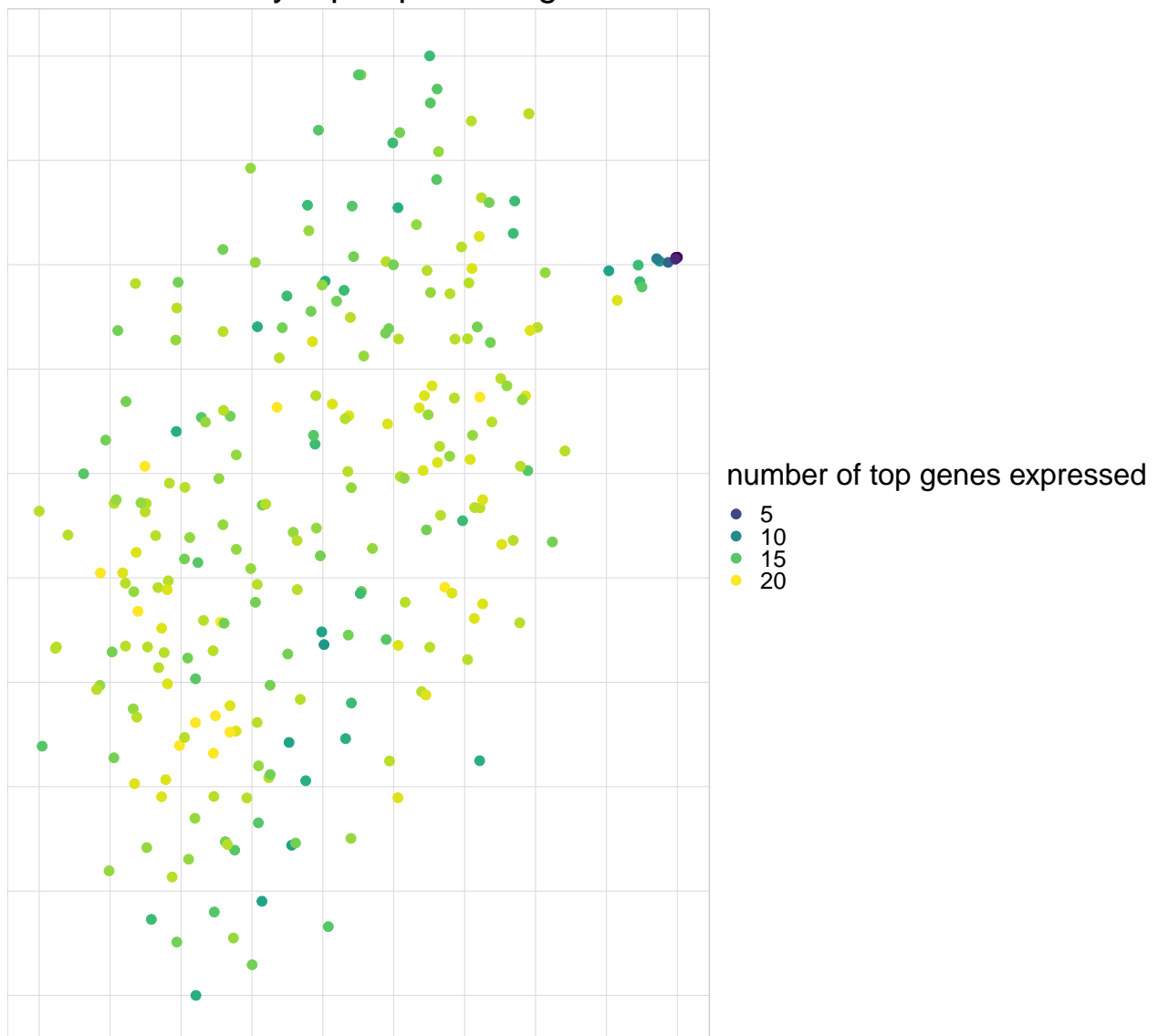


```
[1] "Total number of cells: 660"
```

```
[1] "Top 20 expressing genes: "
```

```
[1] "GAPDH" "IFIT3" "CCR6" "CTLA4" "CCR7" "IL25" "IFIT1"  
[8] "CDC42" "CCR3" "CXCL10" "CXCR4" "CD40" "IRF9" "IFI44"  
[15] "NLRP3" "CXCR3" "BCL2" "SOCS3" "CD28" "JAK2"
```

t-SNE colored by top expressed genes



[1] "Total number of cells: 249"