

# oncoEnrichR Analysis

Mikhail Dozmorov

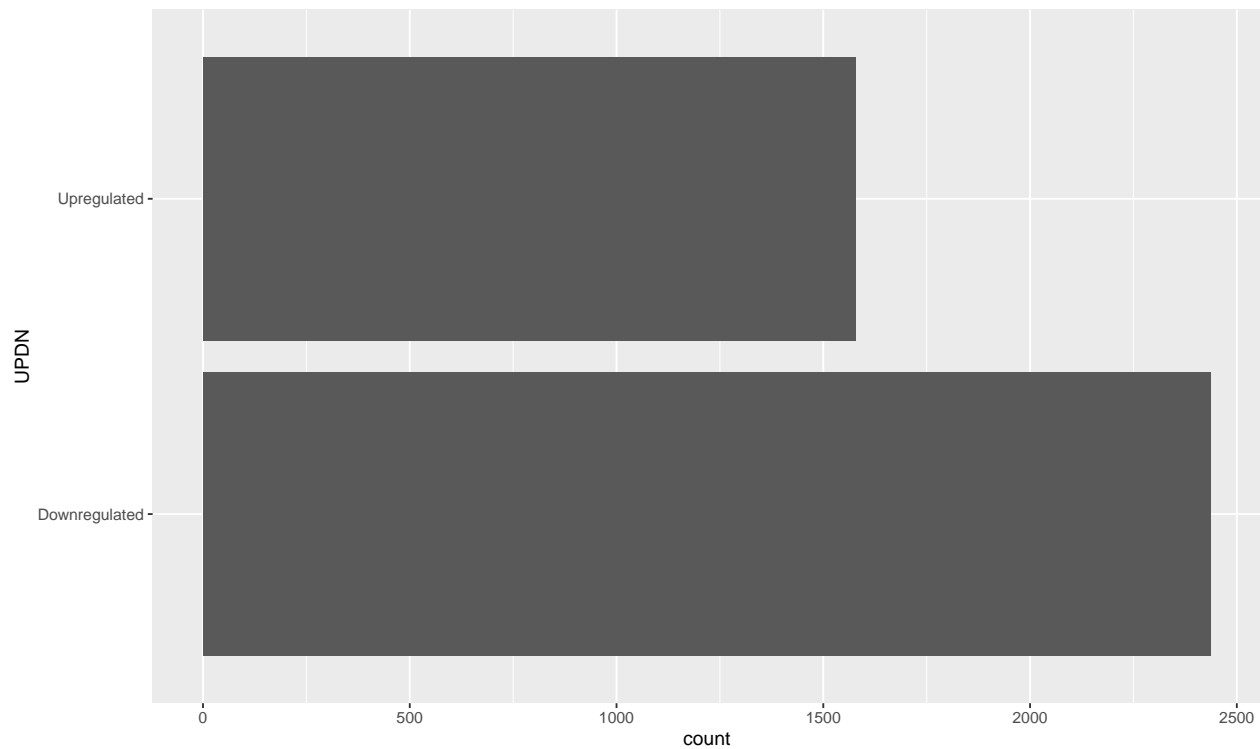
2021-05-12

See <https://github.com/sigven/oncoEnrichR> for an overview of the functionality

## Libraries

## Load data

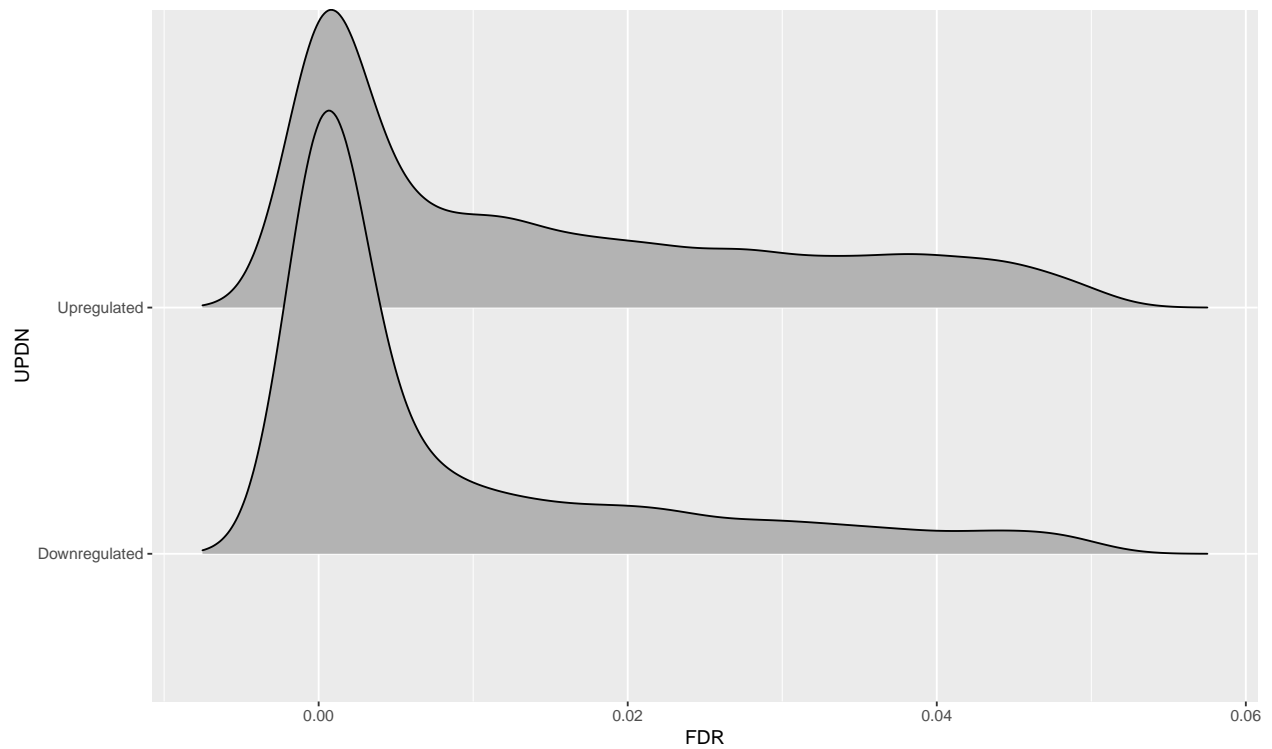
```
[1] "Number of up- and downregulated genes"
```



```
[1] "Summary of the adjusted p-value"
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.0000000	0.0001296	0.0035171	0.0110285	0.0189240	0.0499937

```
[1] "Adjusted p-value discription separately for up- and downregulated genes"
```



## Analysis

```
[1] "Done. Results are in"

[1] "/Users/mdozmorov/Documents/Work/GitHub/RNA-seq/results/oncoEnrichR_SKNSHMBvsSKNSHMA_UP.html"

[1] "/Users/mdozmorov/Documents/Work/GitHub/RNA-seq/results/oncoEnrichR_SKNSHMBvsSKNSHMA._UP.xlsx"

[1] "/Users/mdozmorov/Documents/Work/GitHub/RNA-seq/results/oncoEnrichR_SKNSHMBvsSKNSHMA_DN.html"

[1] "/Users/mdozmorov/Documents/Work/GitHub/RNA-seq/results/oncoEnrichR_SKNSHMBvsSKNSHMA._DN.xlsx"
```