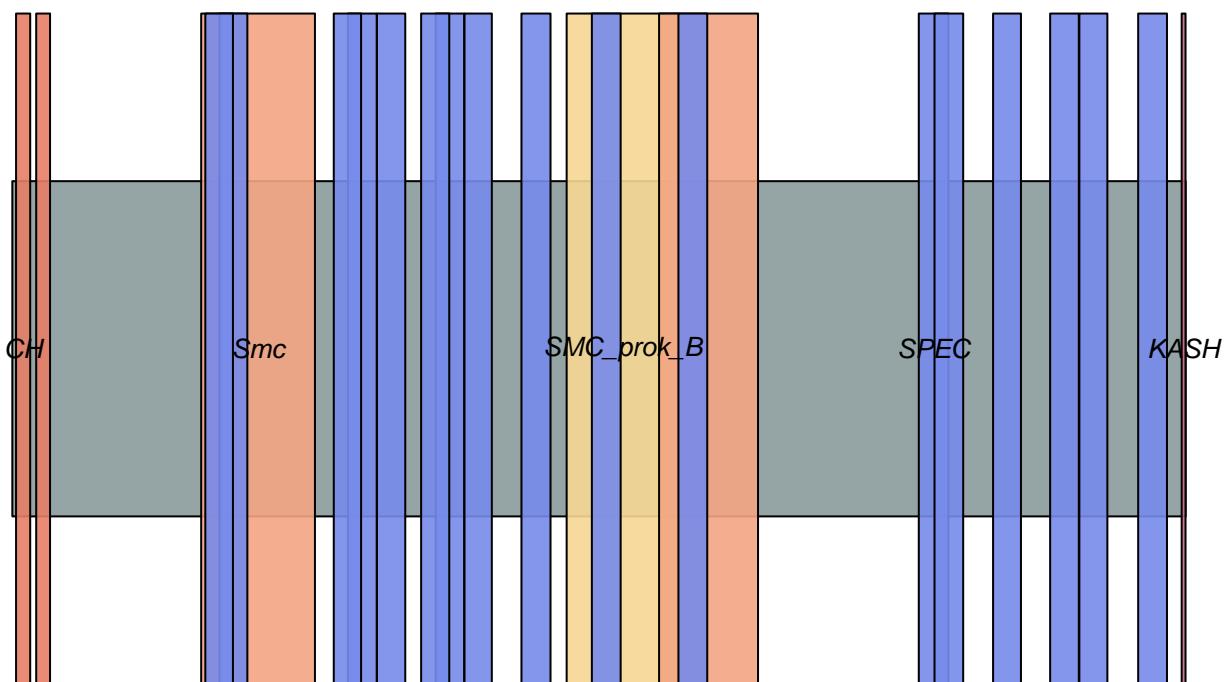


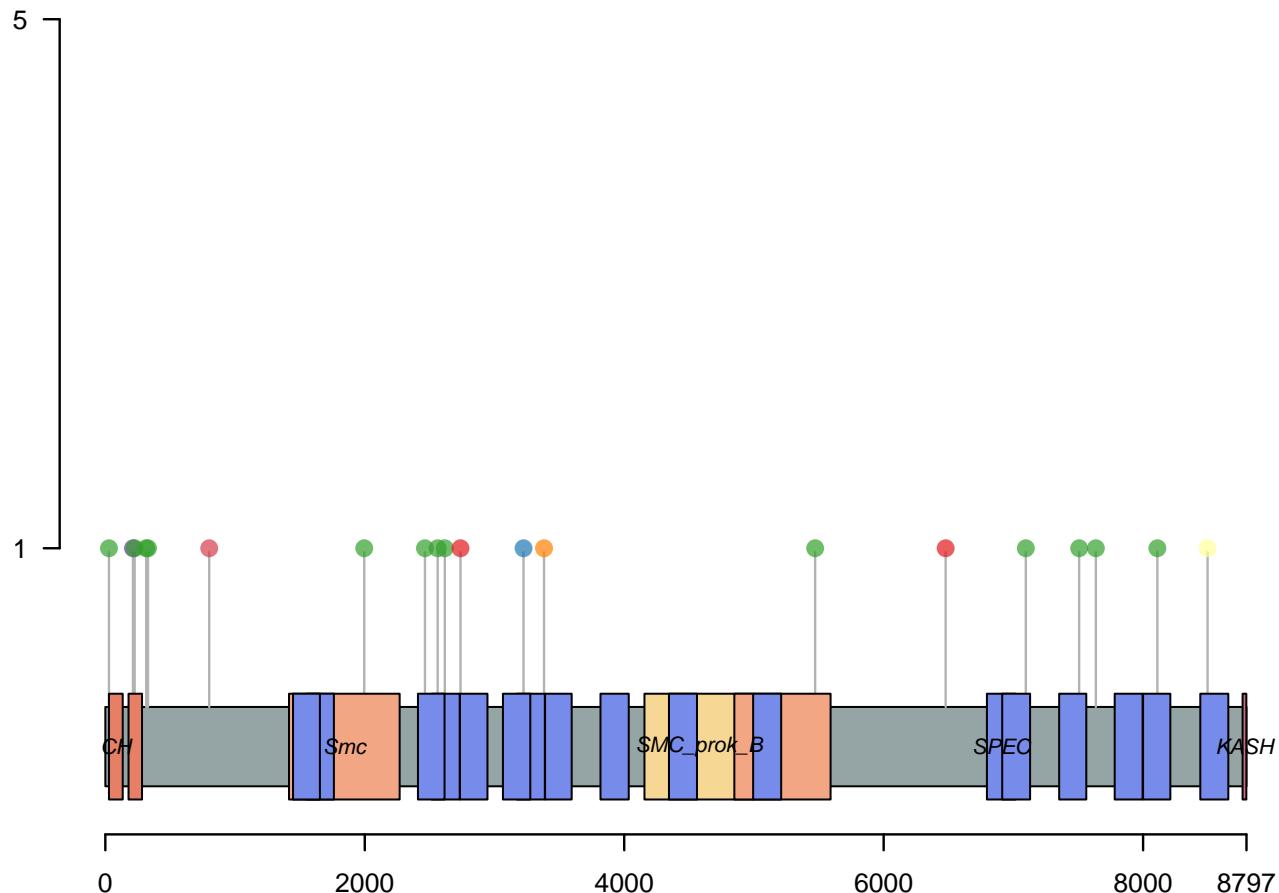
*SYNE1*

NM\_182961



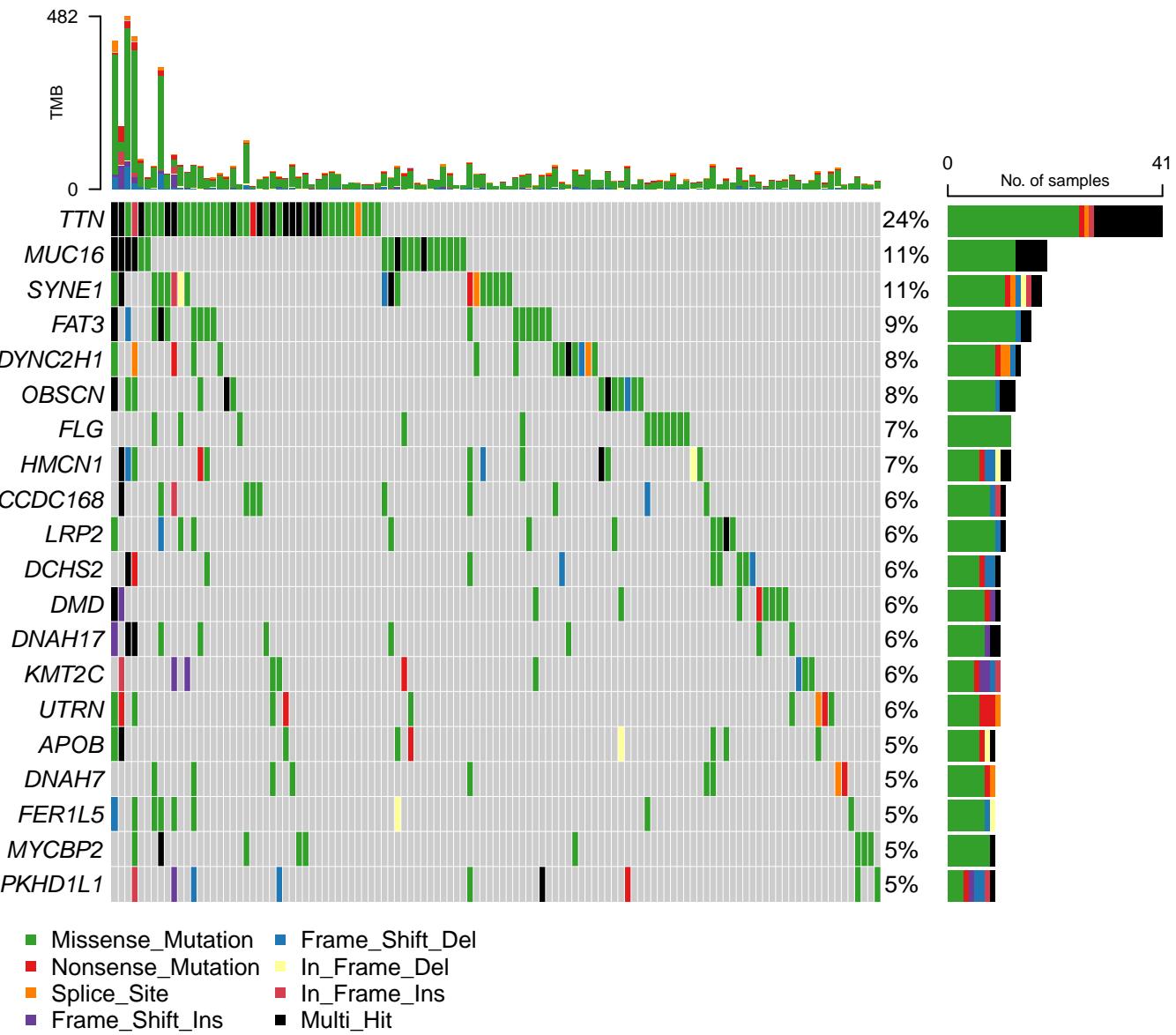
# SYNE1 : [Somatic Mutation Rate: 10.53%]

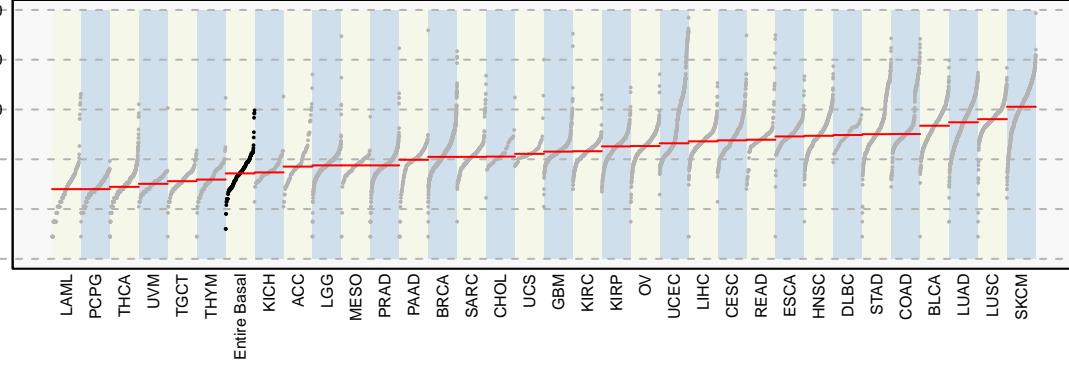
NM\_182961



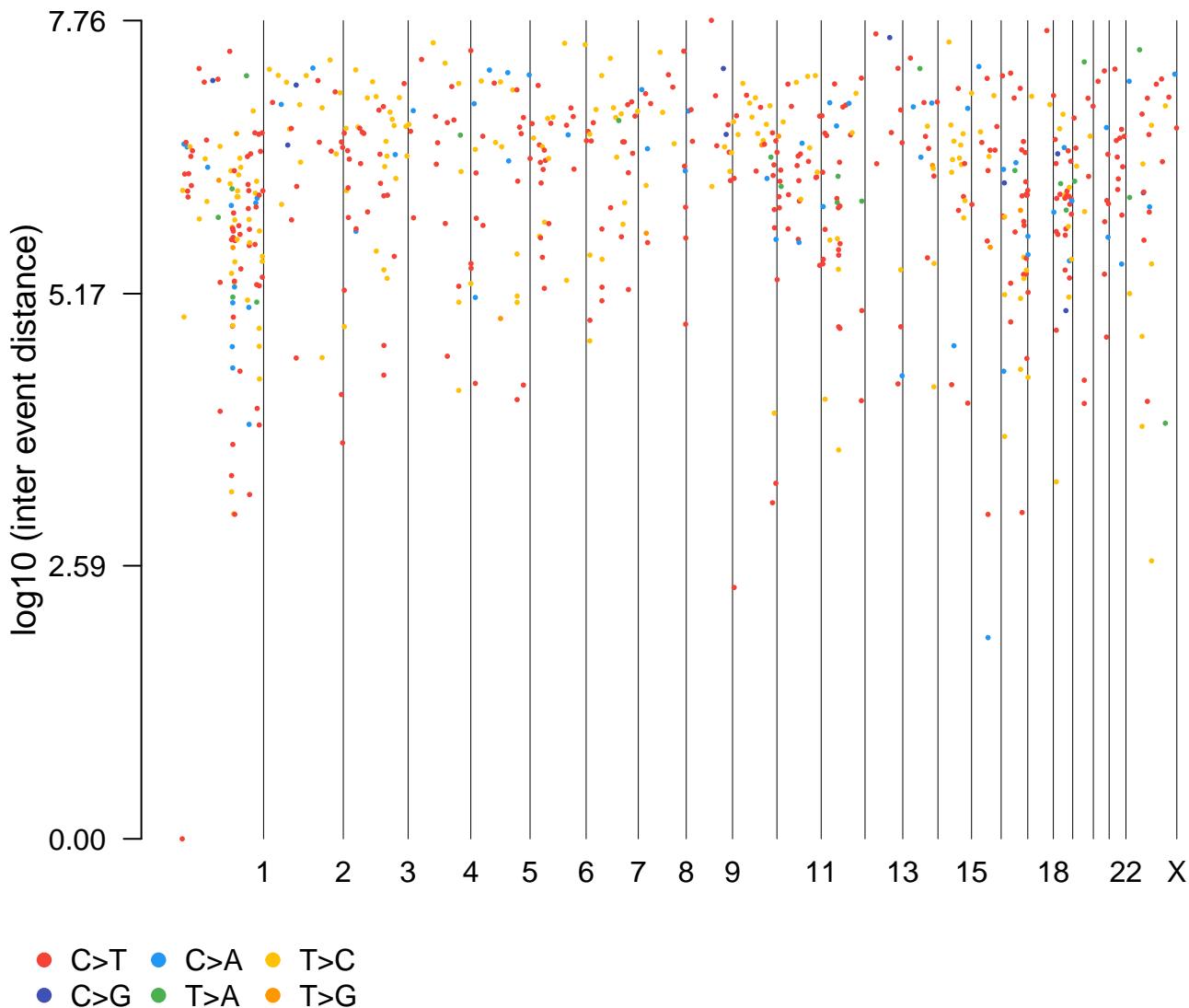
- Missense\_Mutation
- Frame\_Shift\_Del
- Frame\_Shift\_Ins
- Splice\_Site
- In\_Frame\_Ins
- In\_Frame\_Del
- Nonsense\_Mutation

# Altered in 117 (68.42%) of 171 samples.

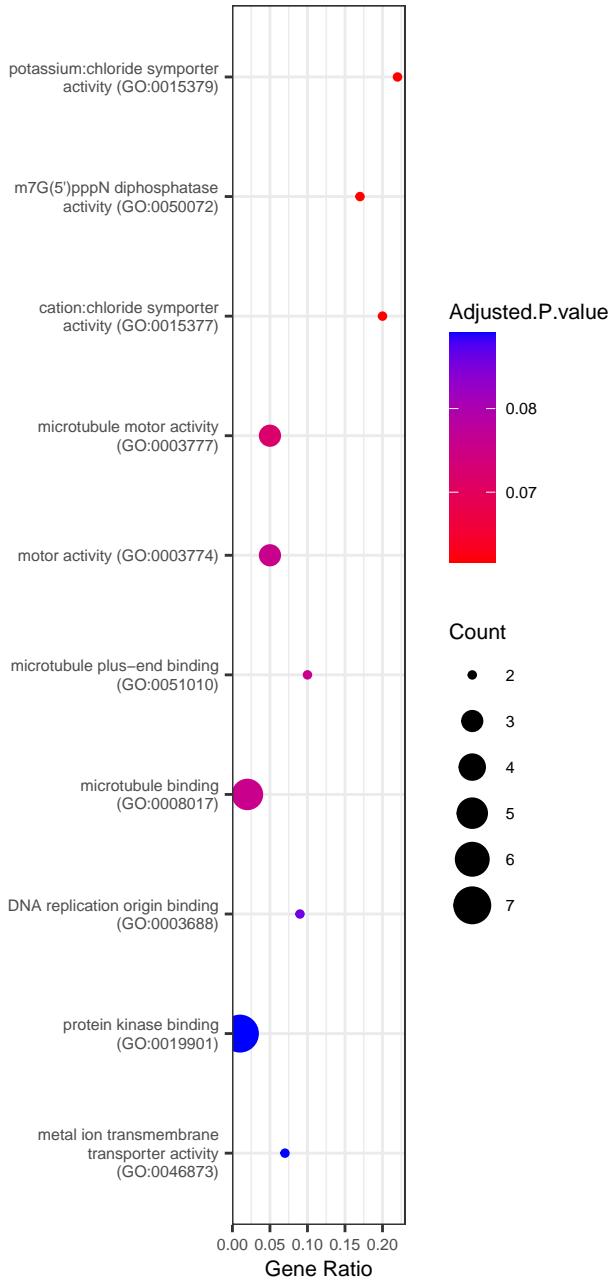




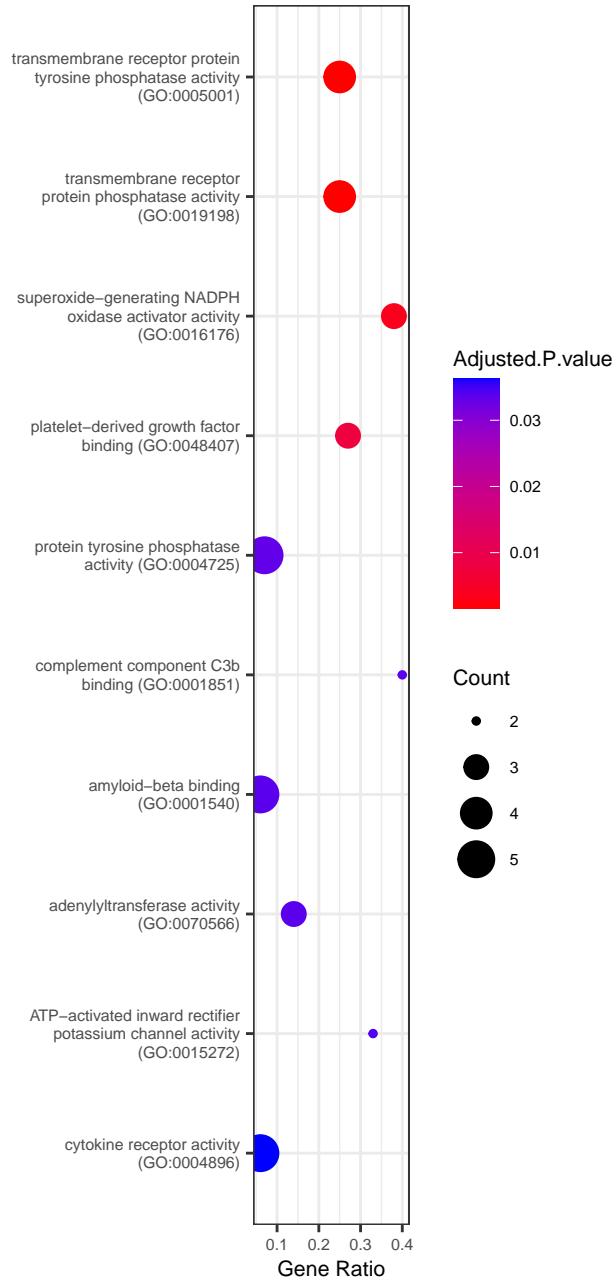
# TCGA-BH-A18G-01A-11D-A12B-09



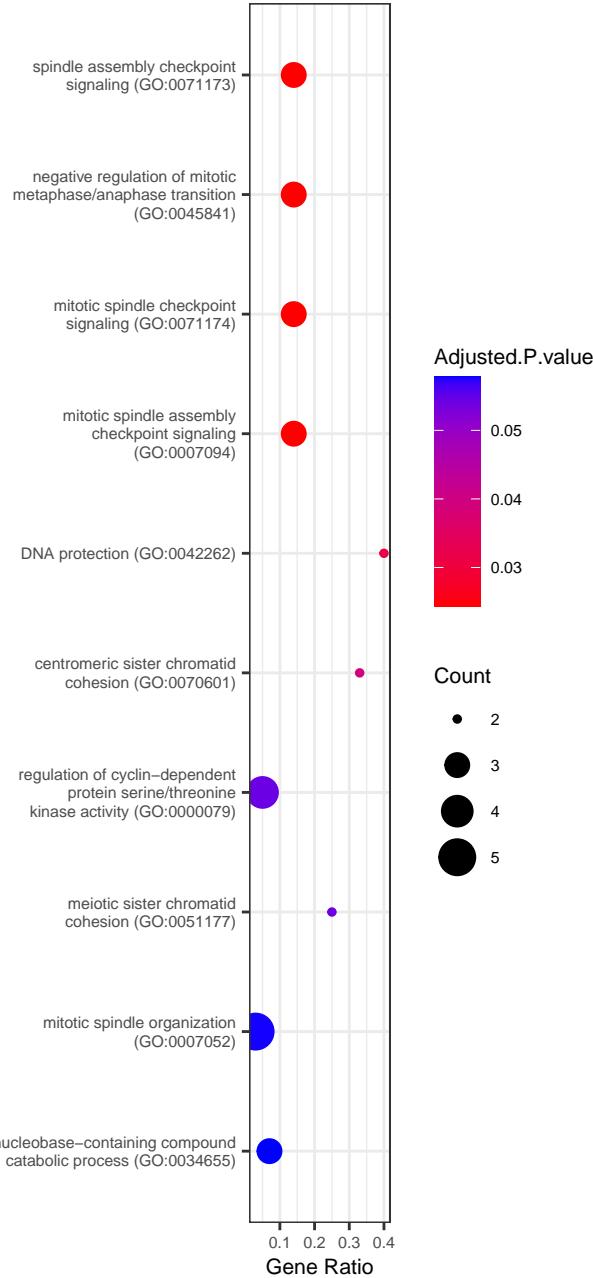
## Oncogenes



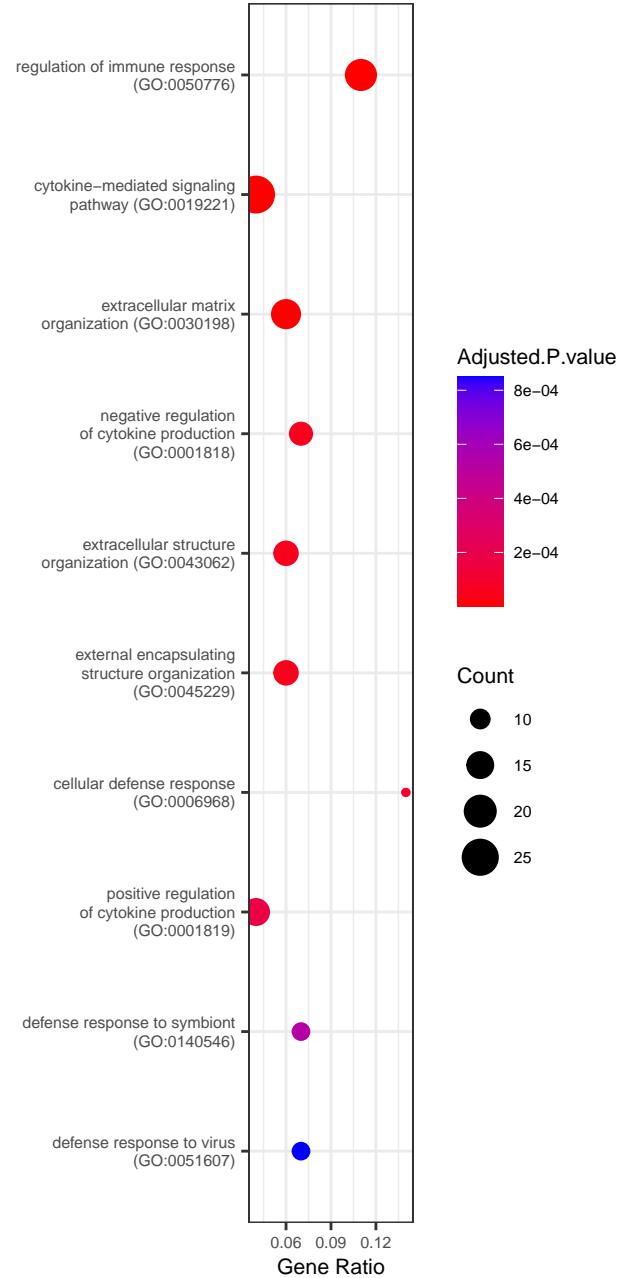
## Tumor suppressors

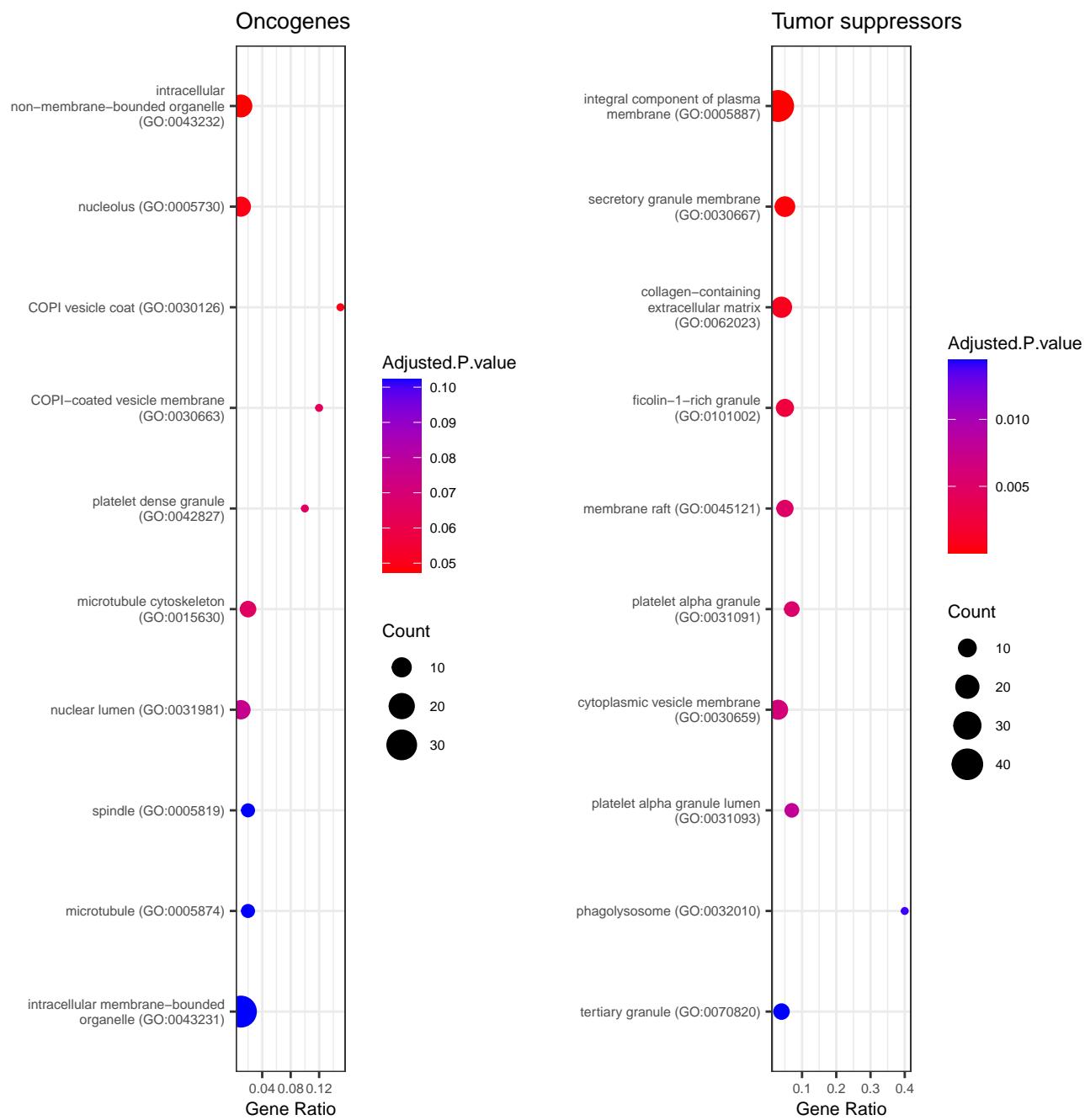


## Oncogenes

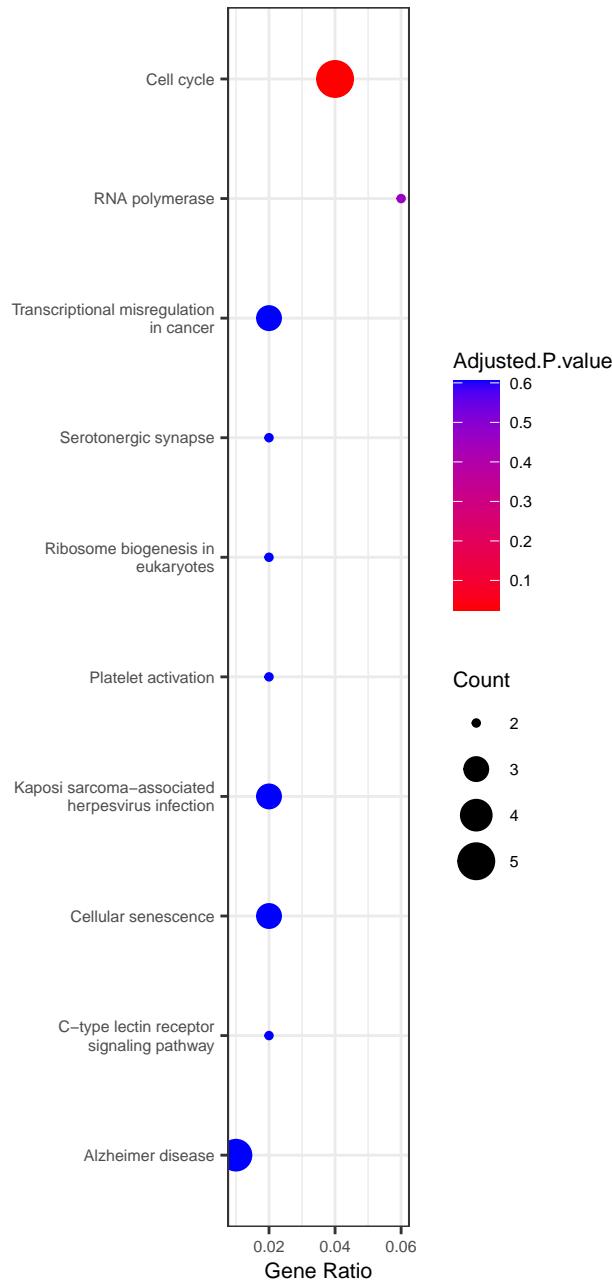


## Tumor suppressors





## Oncogenes



## Tumor suppressors

