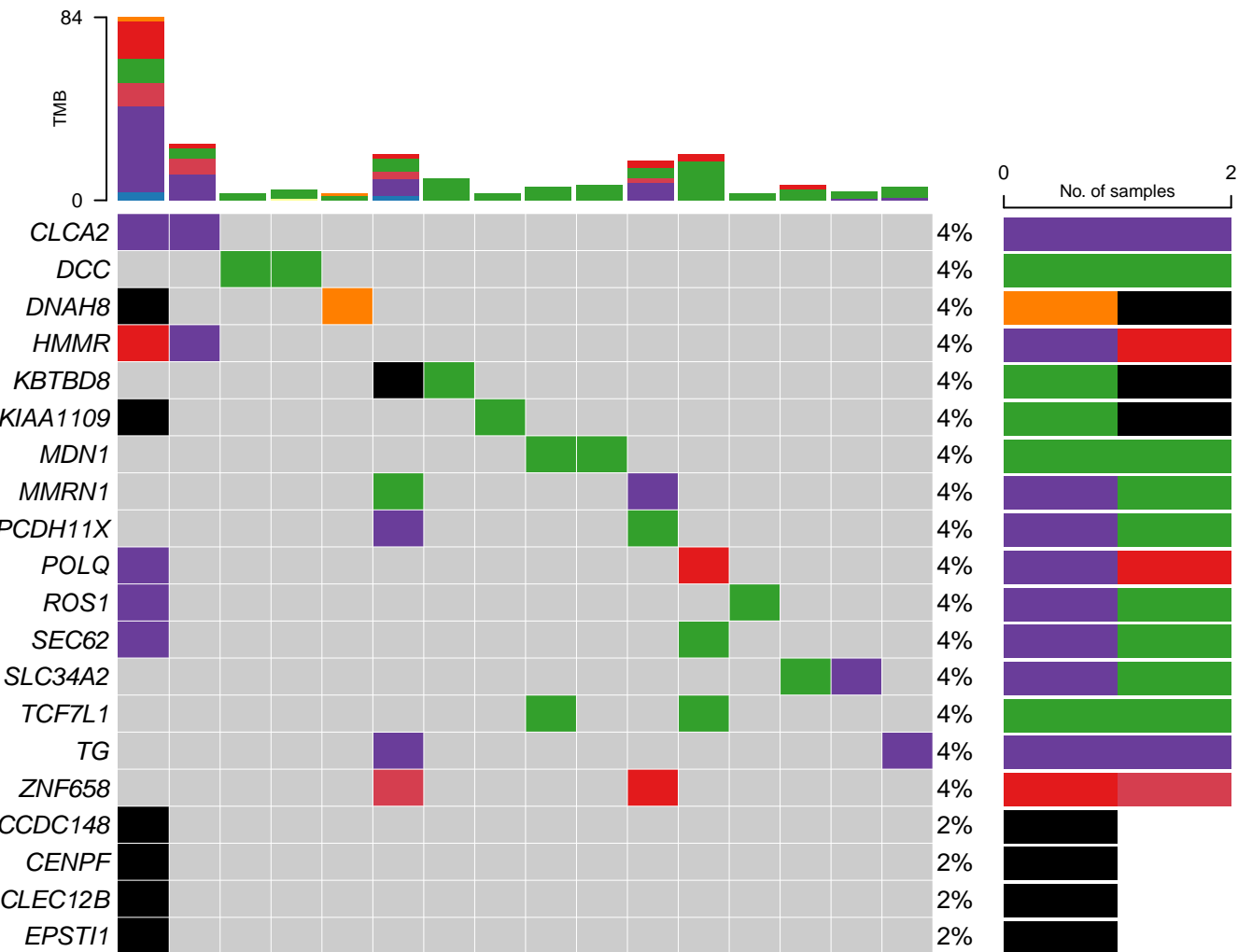
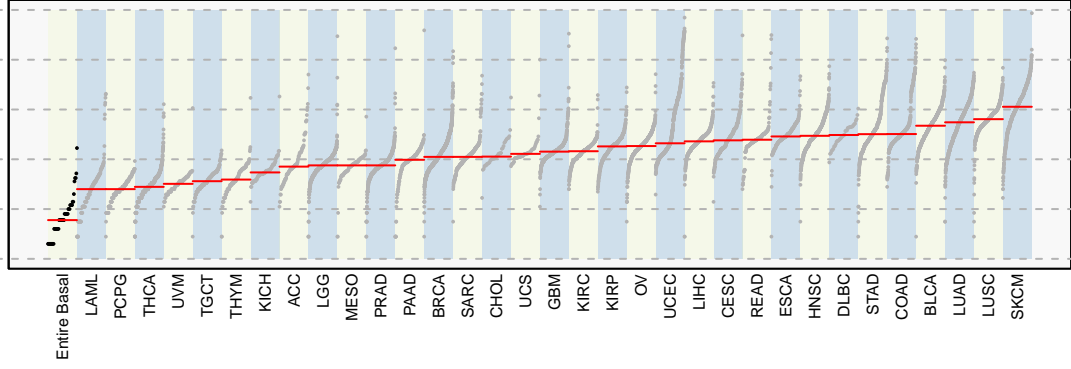


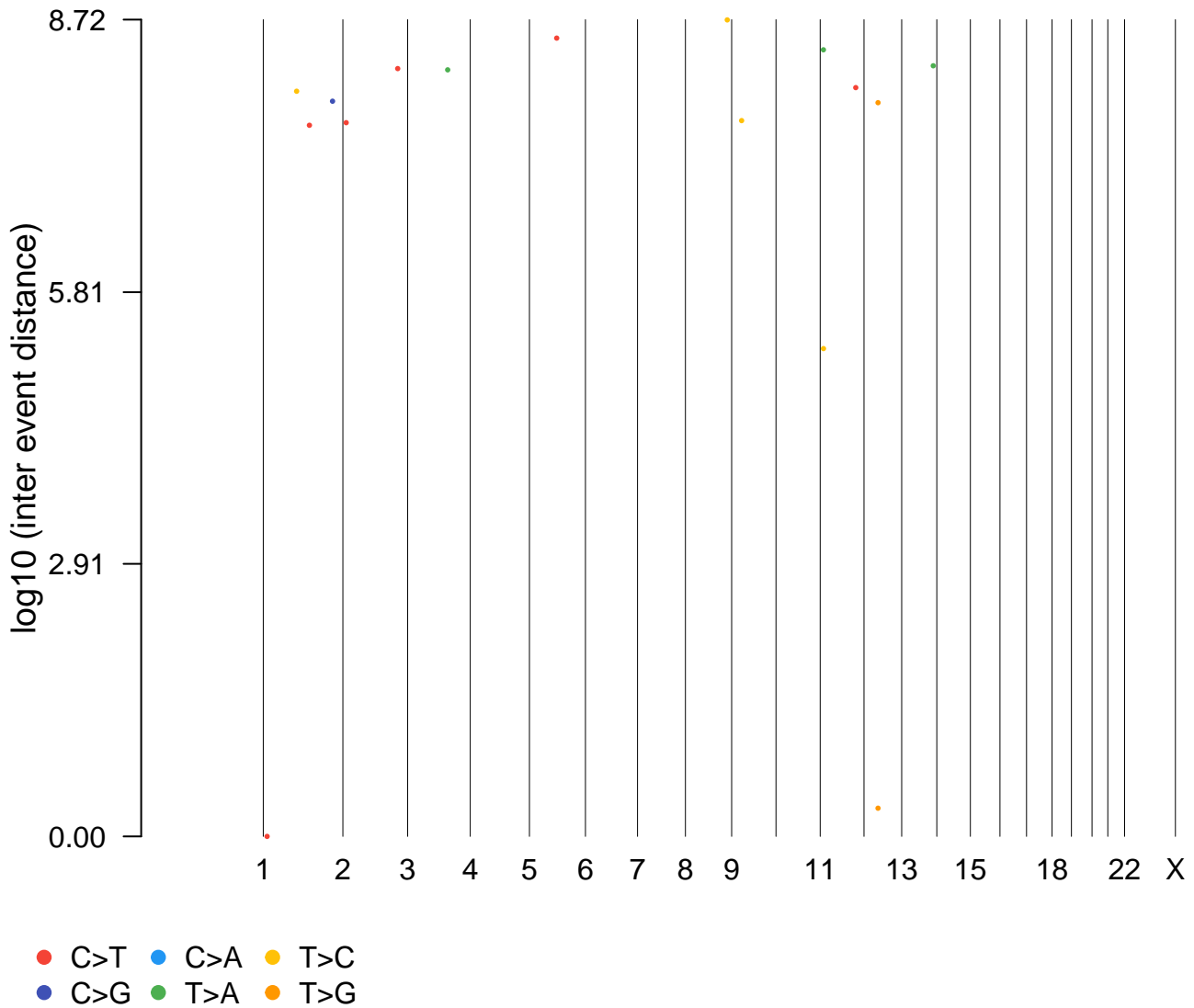
Altered in 16 (30.77%) of 52 samples.



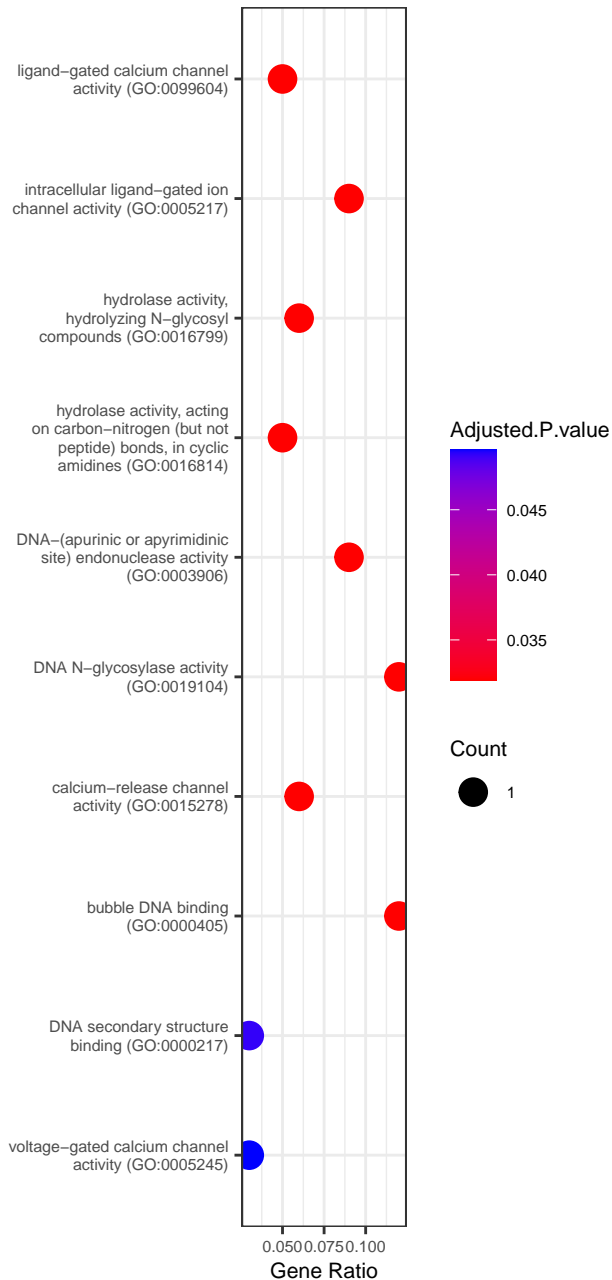
Frame_Shift_Ins
 Missense_Mutation
 Frame_Shift_Del
 Splice_Site
 Nonsense_Mutation
 In_Frame_Ins
 Multi_Hit



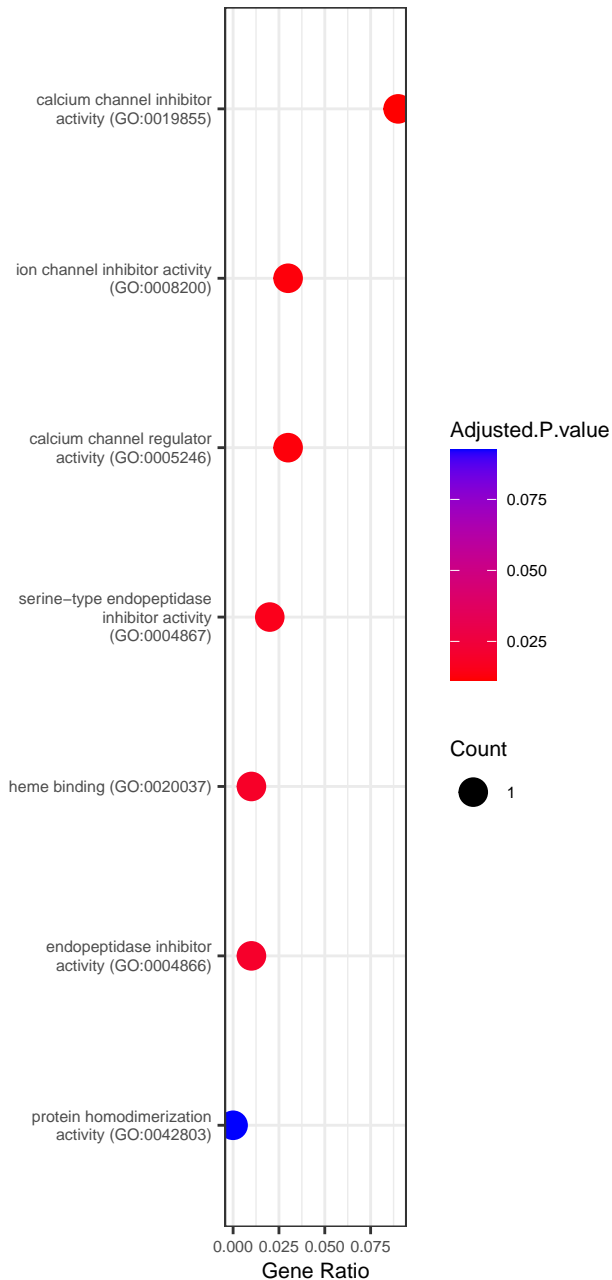
TCGA-EL-A3ZG-01A-11D-A23M-08



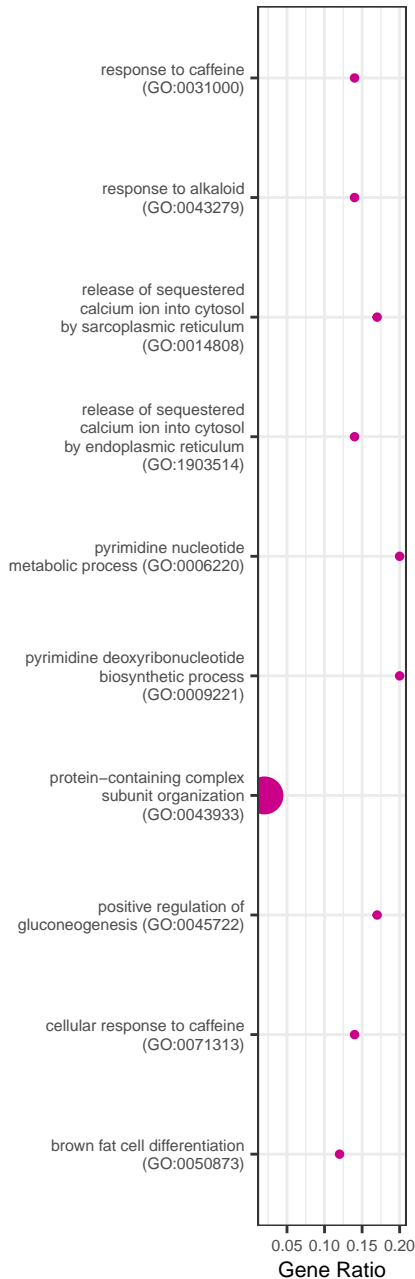
Oncogenes



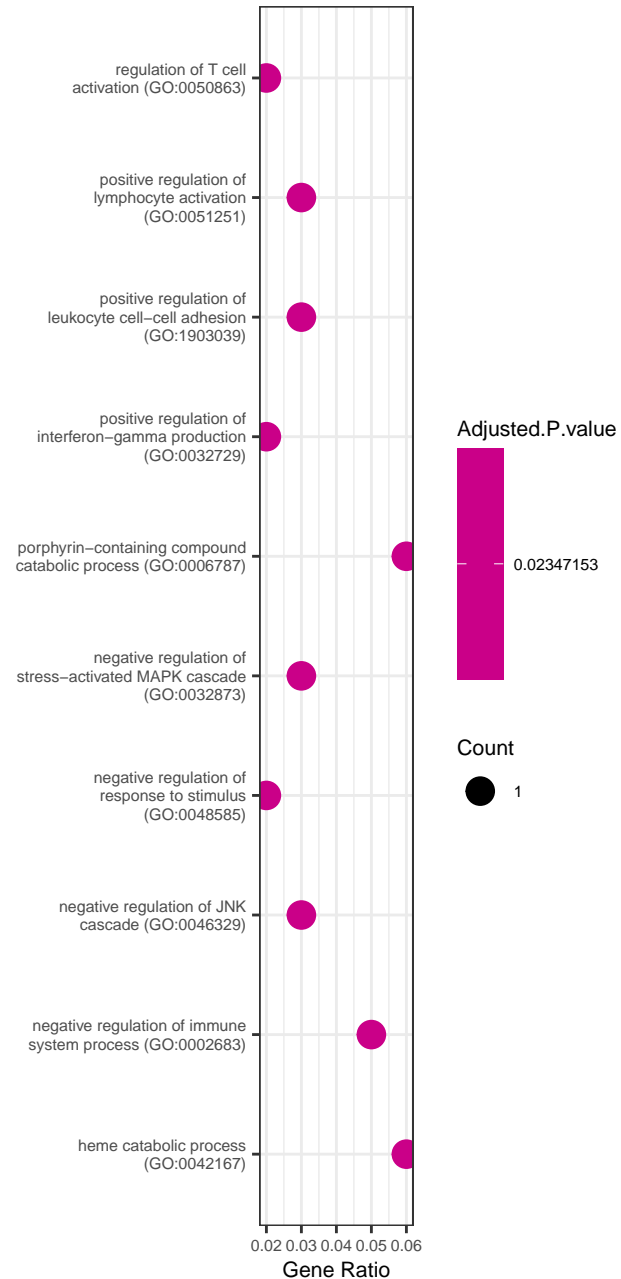
Tumor suppressors



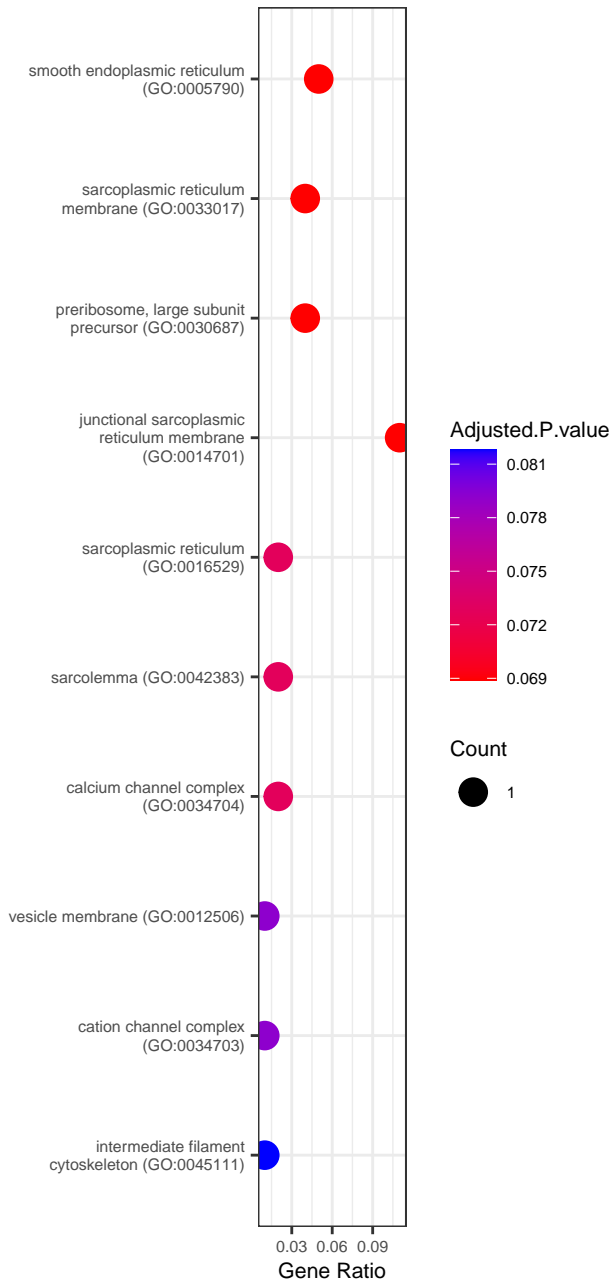
Oncogenes



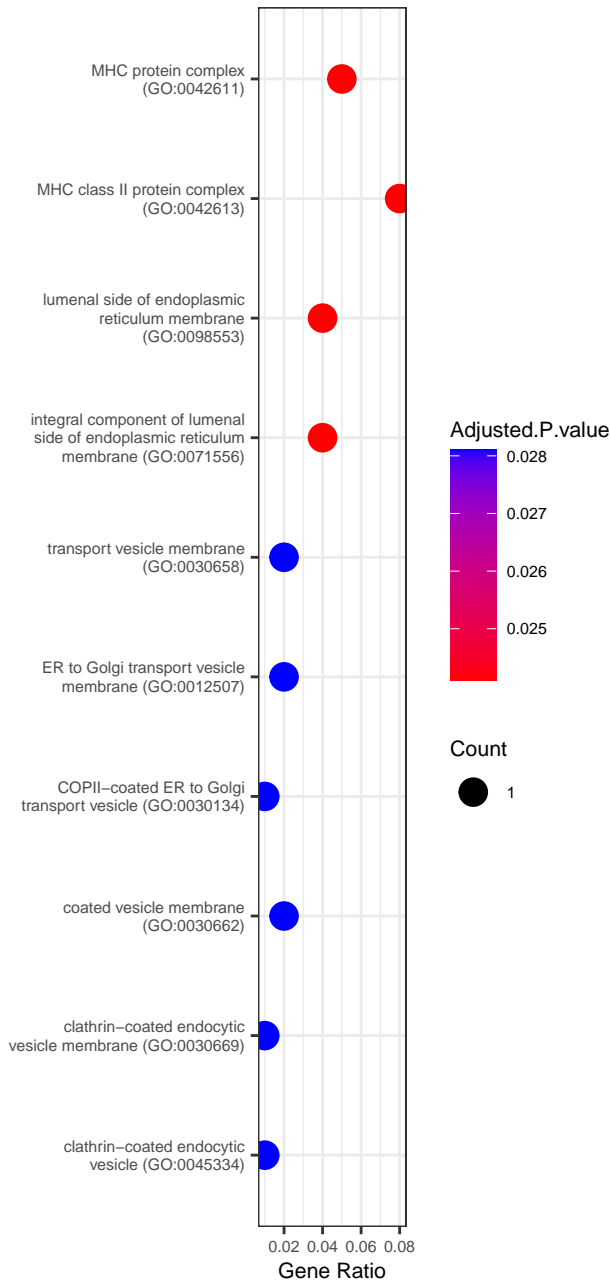
Tumor suppressors



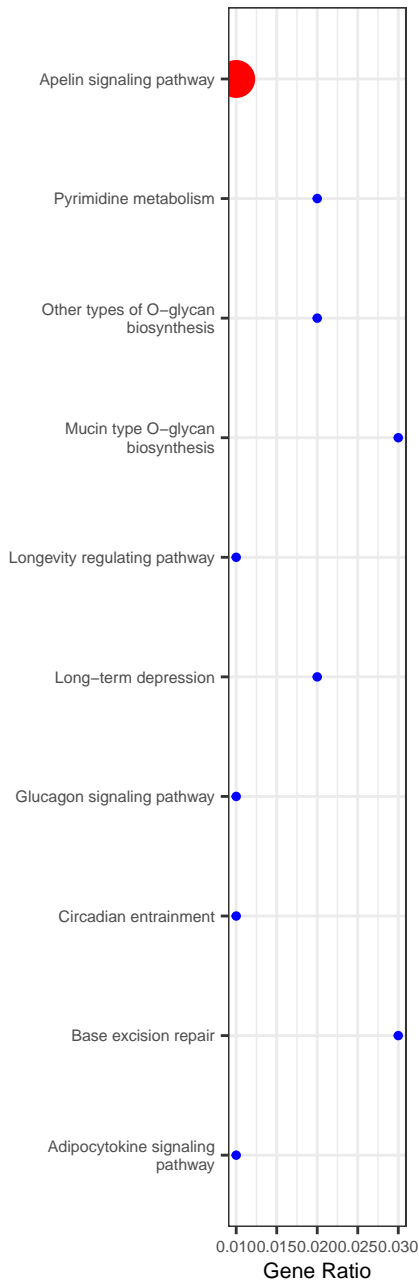
Oncogenes



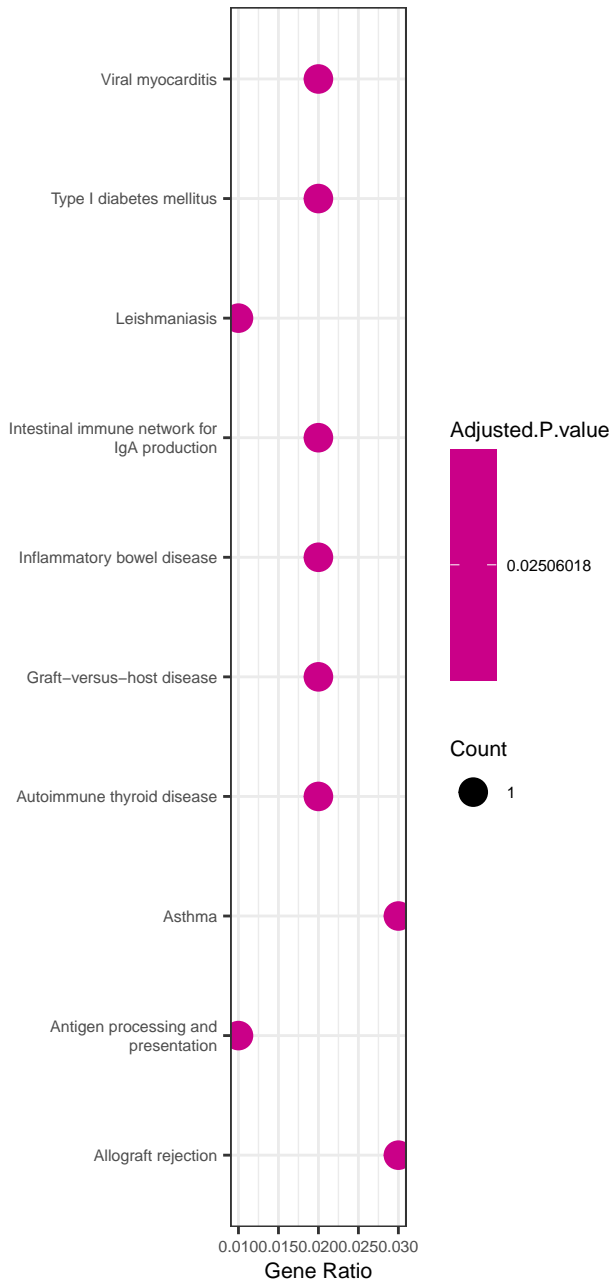
Tumor suppressors



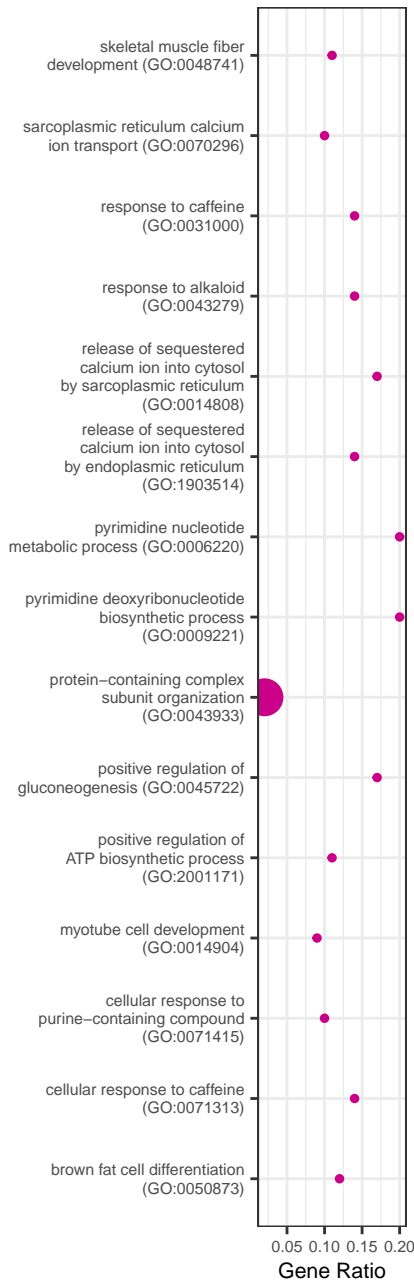
Oncogenes



Tumor suppressors



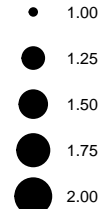
Oncogenes



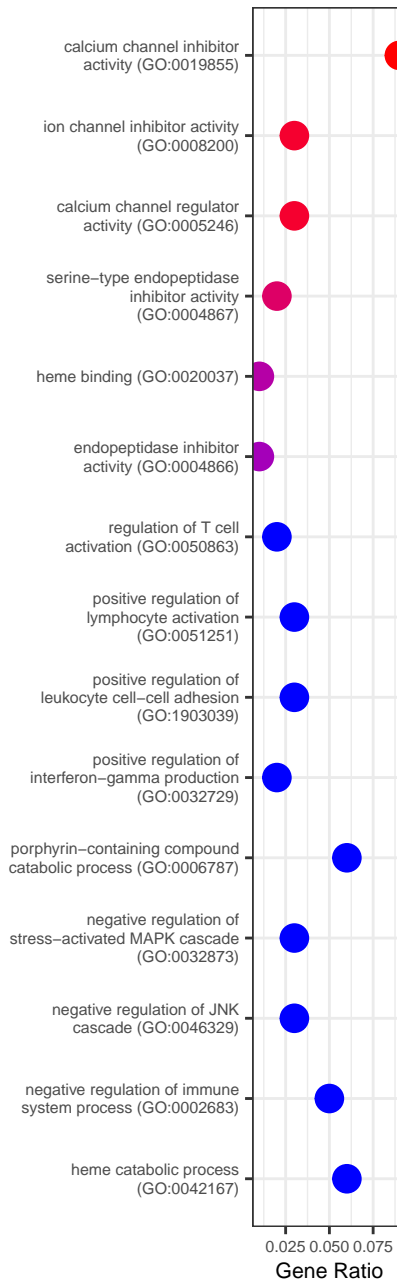
Adjusted.P.value

0.03141782

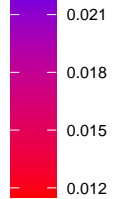
Count



Tumor suppressors



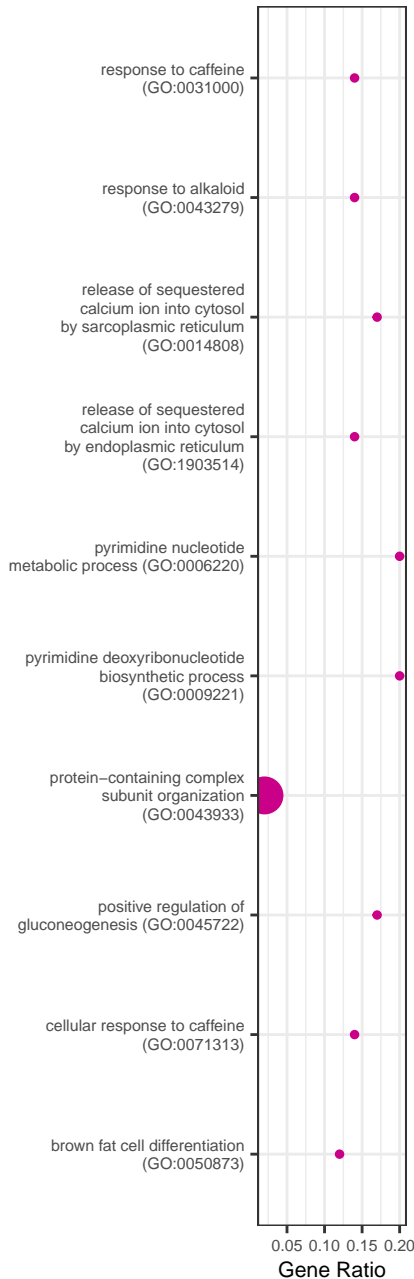
Adjusted.P.value



Count



Oncogenes



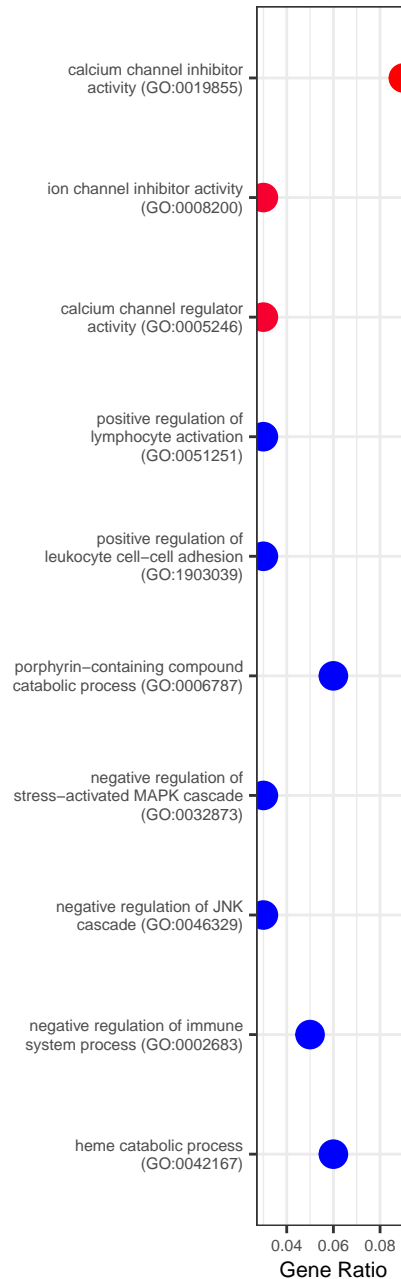
Adjusted.P.value

0.03141782

Count



Tumor suppressors



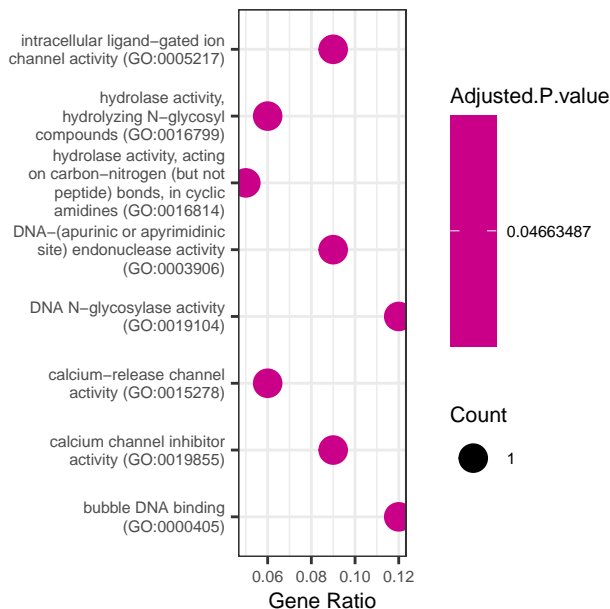
Adjusted.P.value

0.021
0.018
0.015
0.012

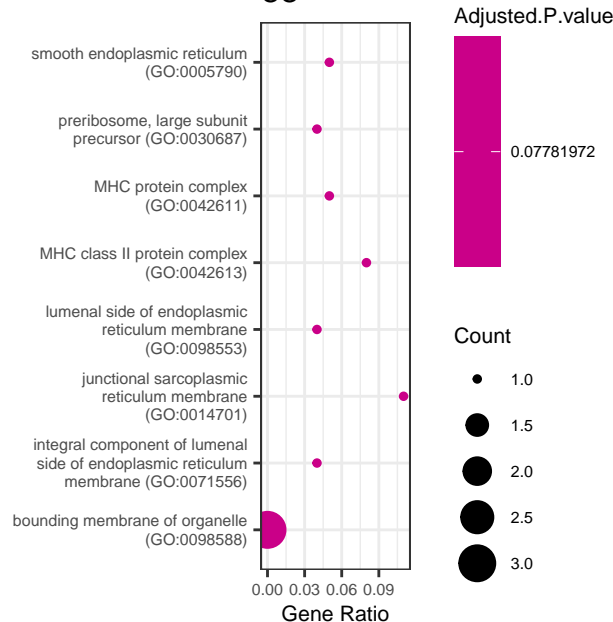
Count

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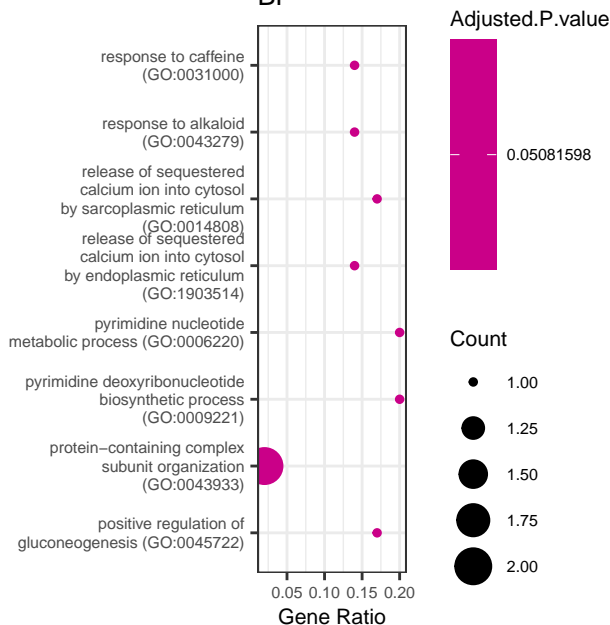
MF



CC



BP



KEGG

