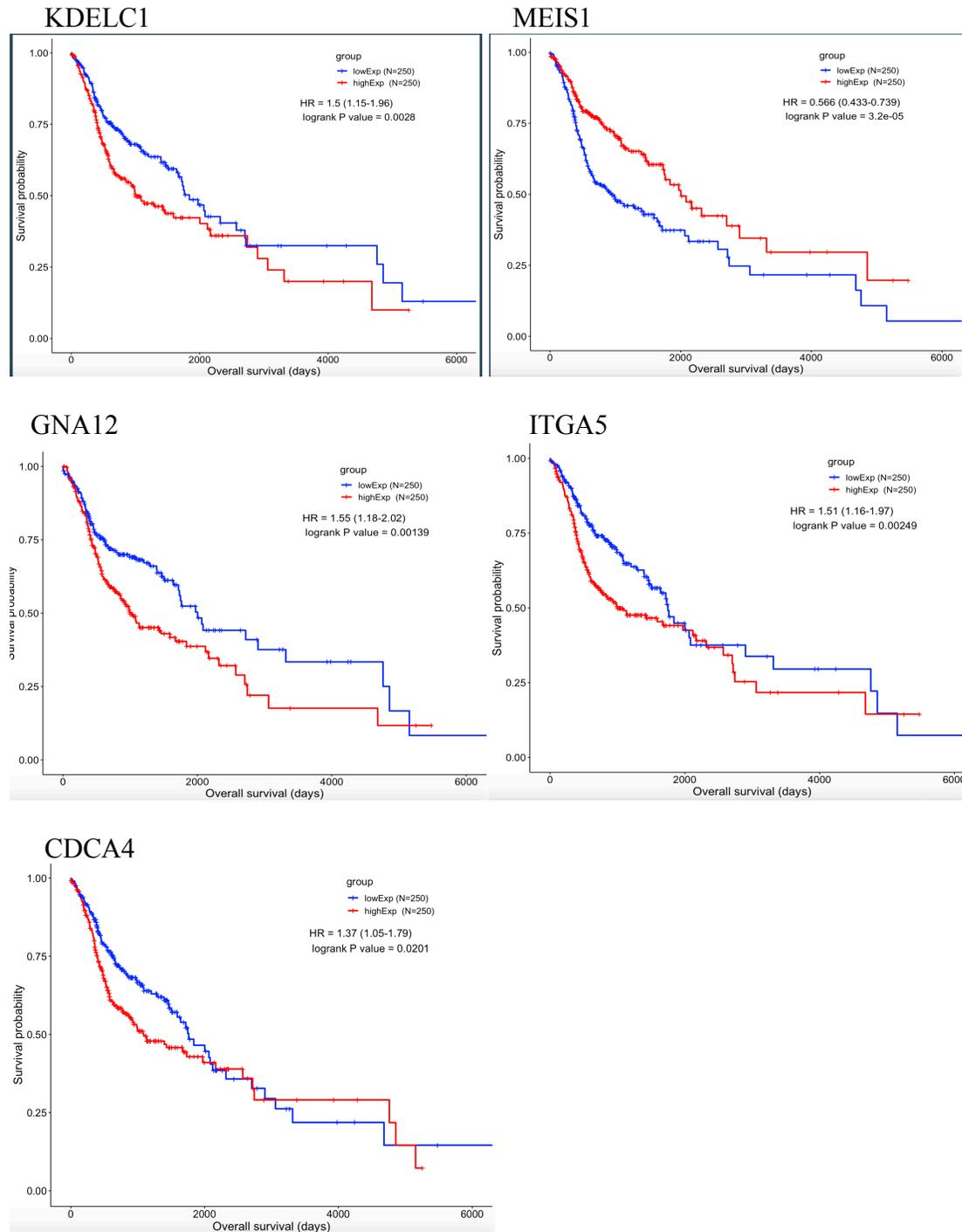
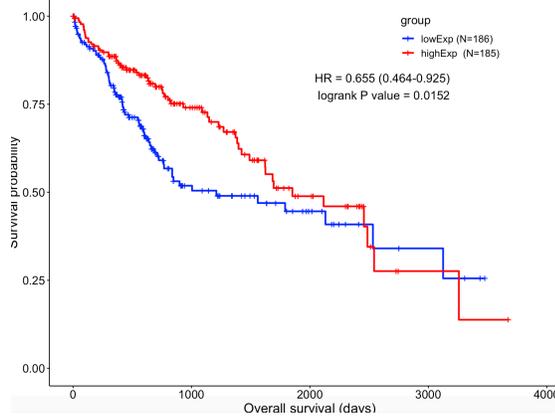
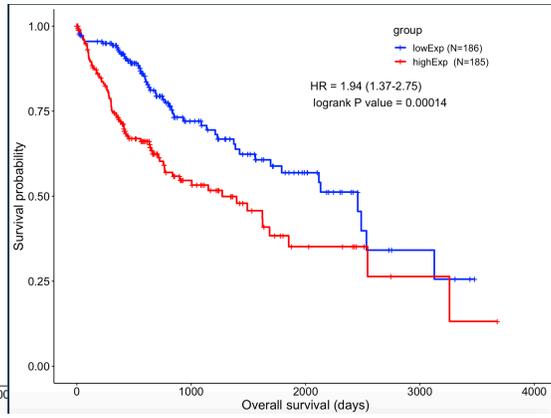
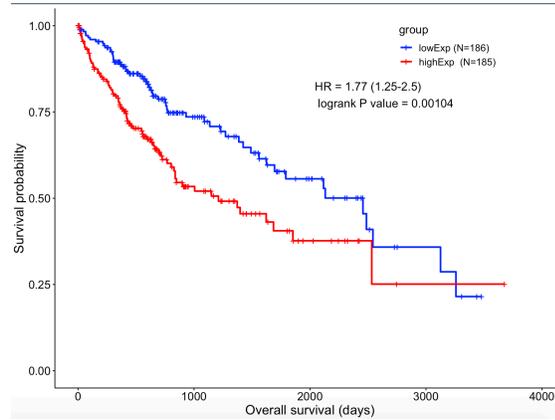
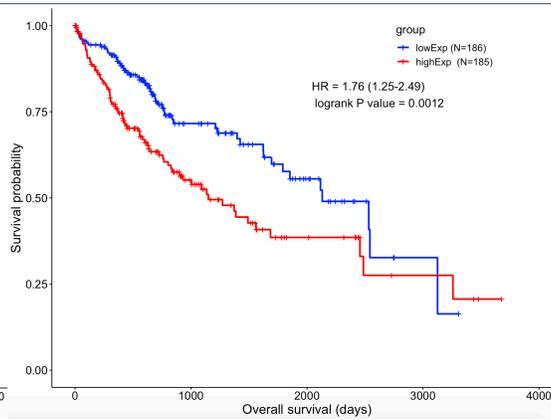
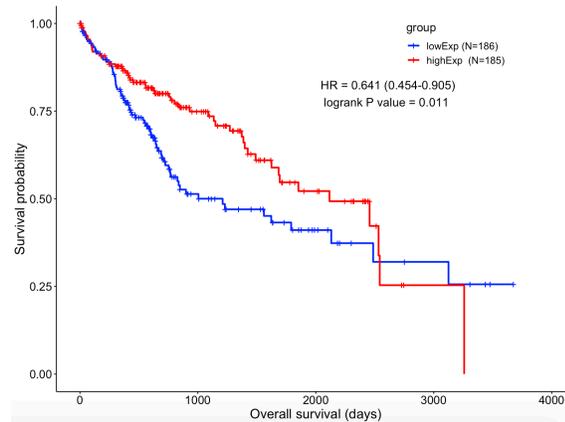
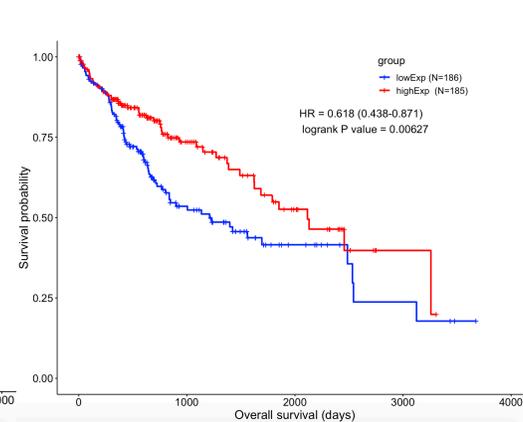


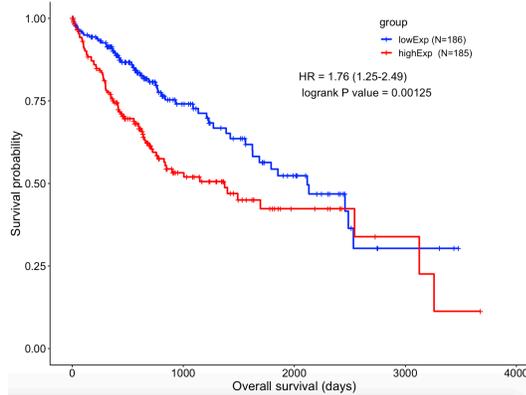
Supplementary Figure 2. OS analysis of genes in patients with TCGA cancers. A: HNSC, B: LIHC, C: READ, D: STAD, E: COAD.

(A) HNSC

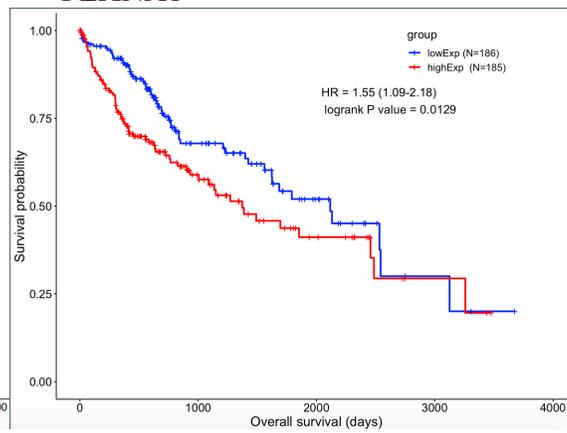


(B) LIHC
RUNX3

CDC7

SCML2

LIMK1

CD69

RIPOR2


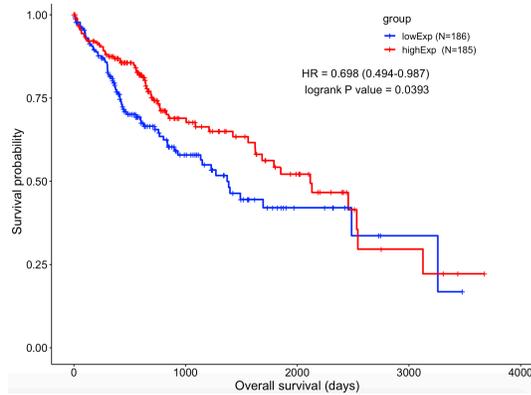
FANCE



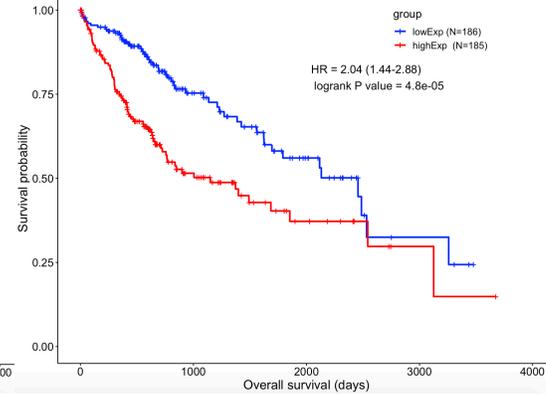
PLXNA1



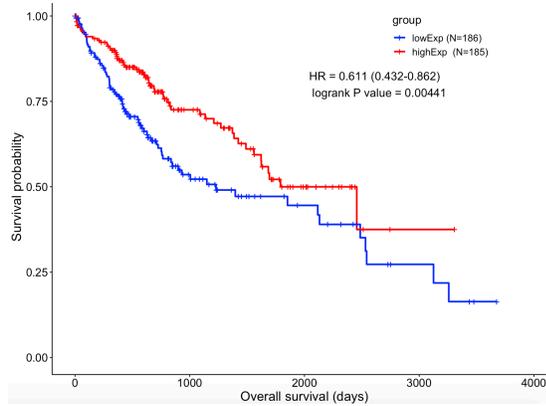
GADD45A



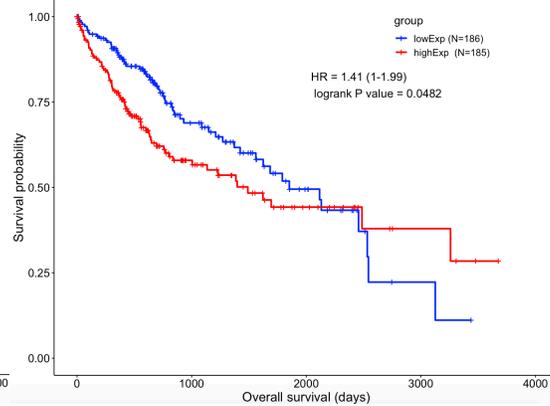
E2F8



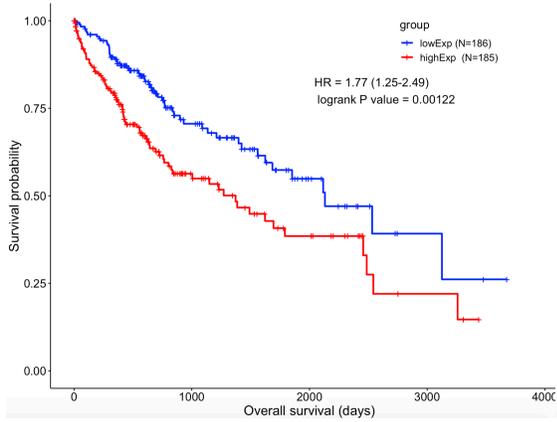
EFNA3



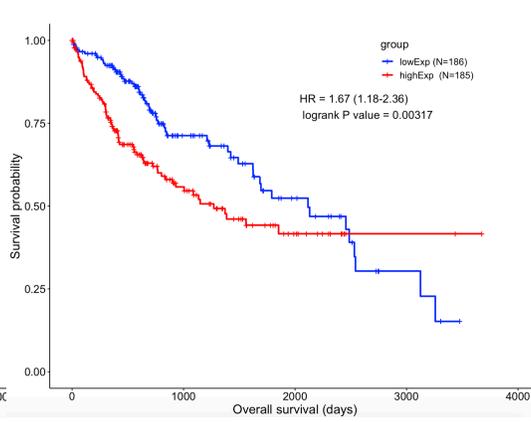
SYBU



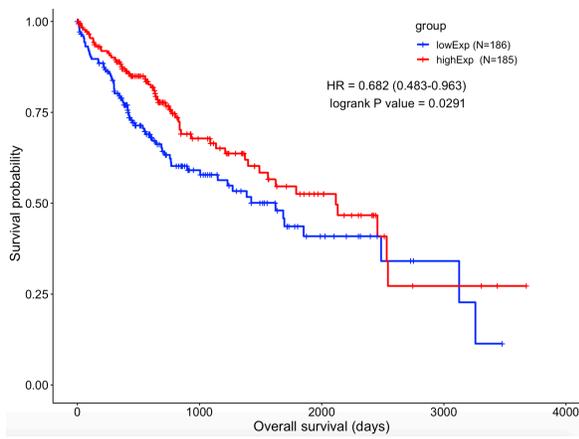
SLC7A11



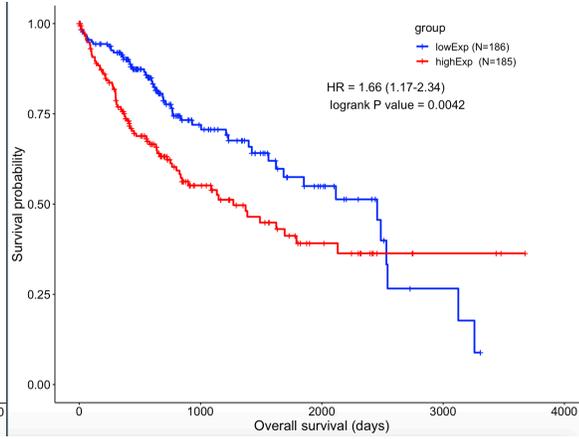
ENAH



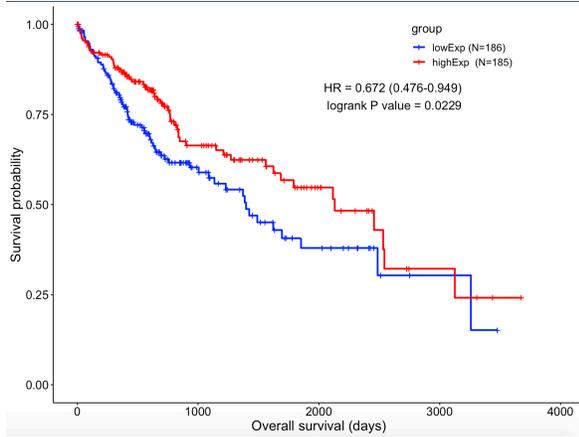
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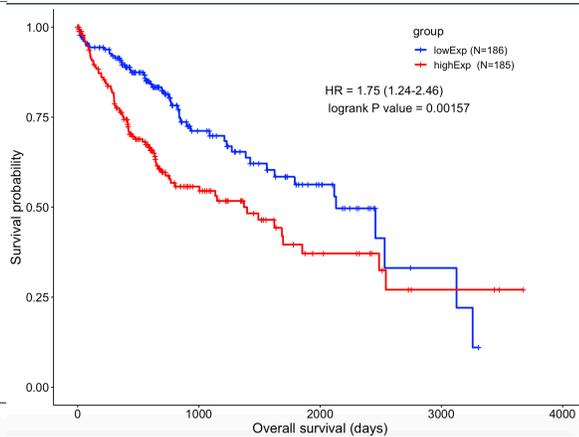
TMEM164

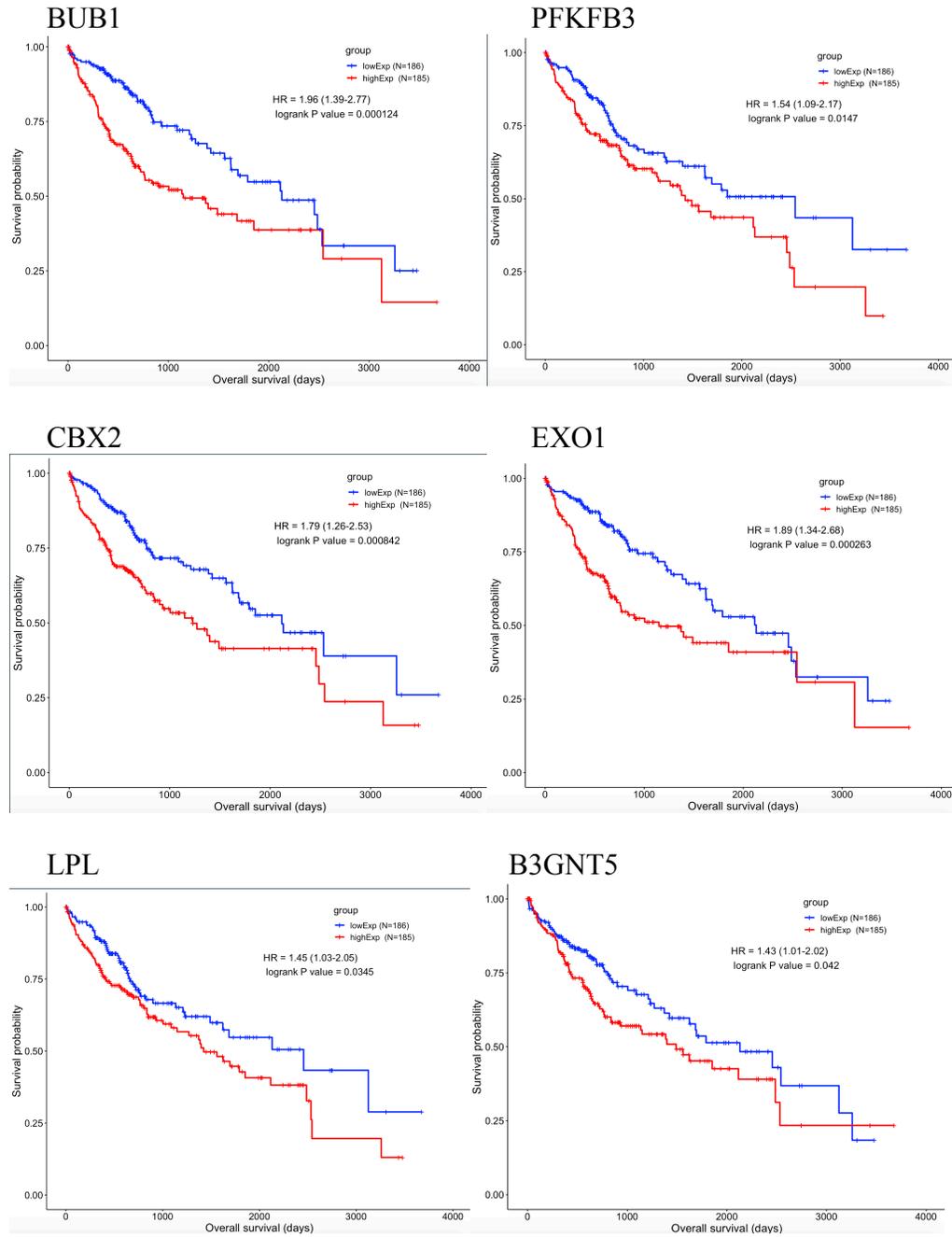


AUTS2

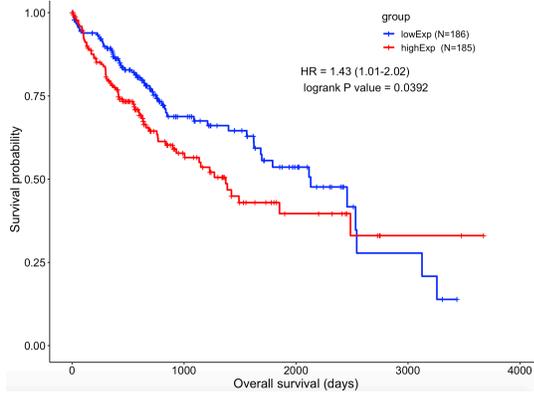


H2AFZ

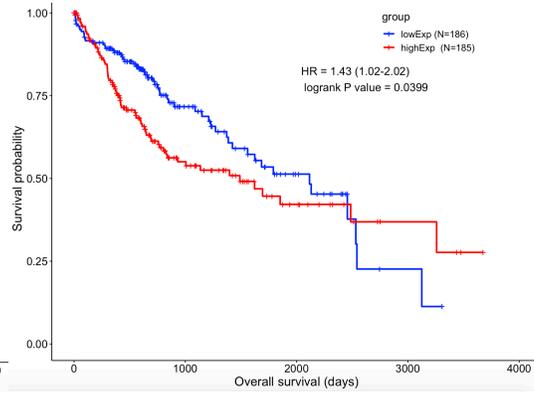




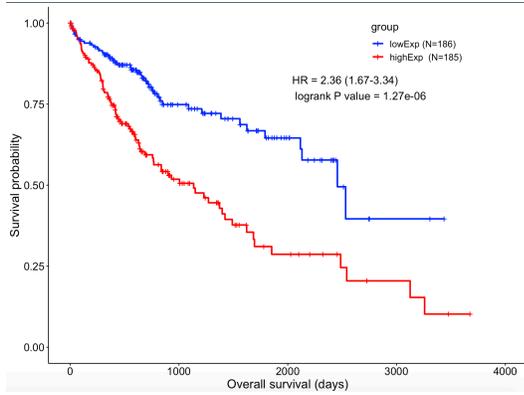
LMNB2



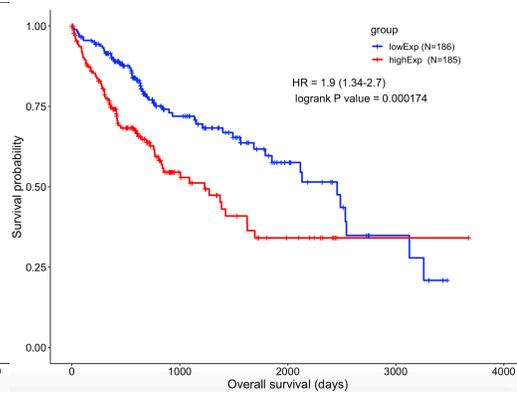
SOX12



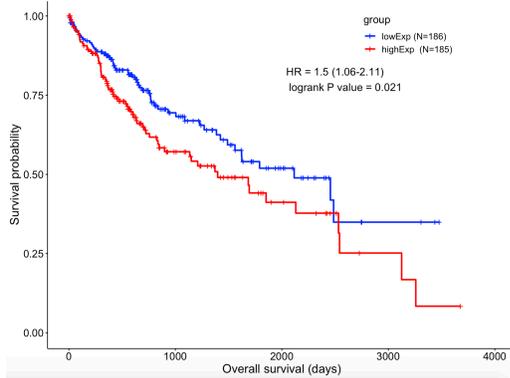
JPT1



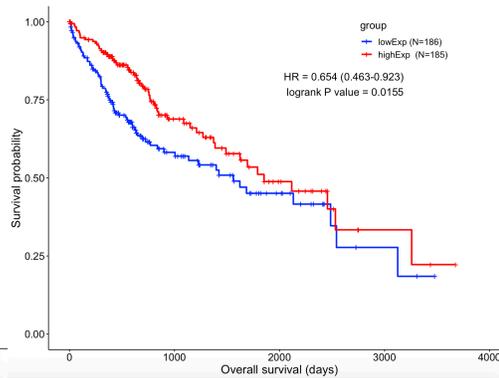
MAFG



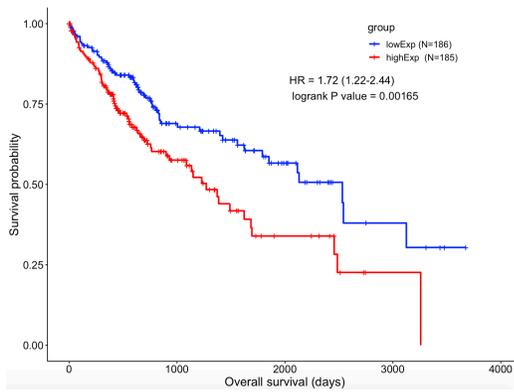
DNM3



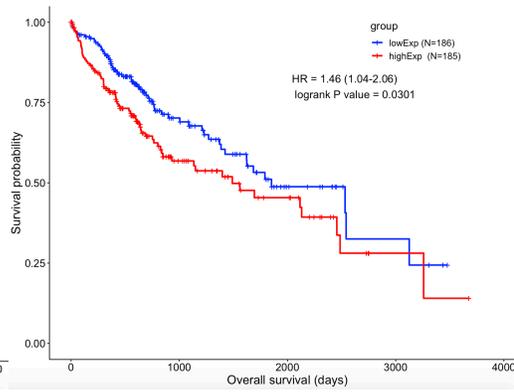
ZNF607



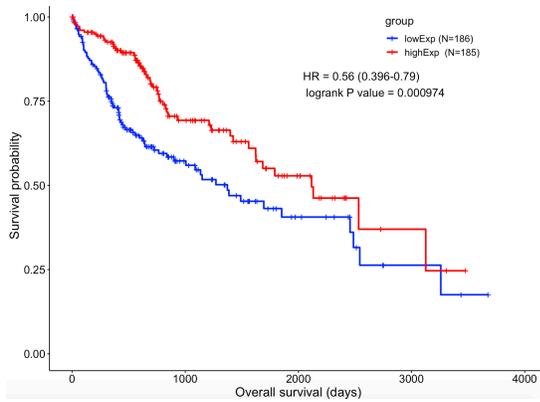
STK39



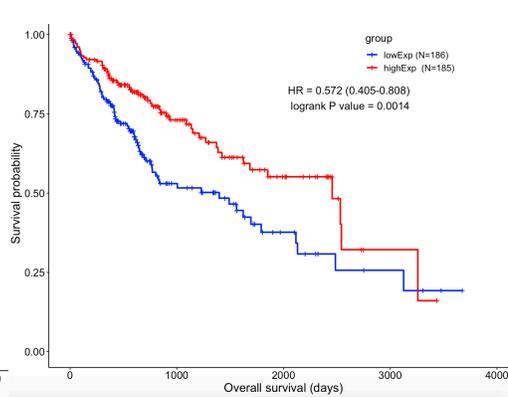
ANKRD13B



SMOC1

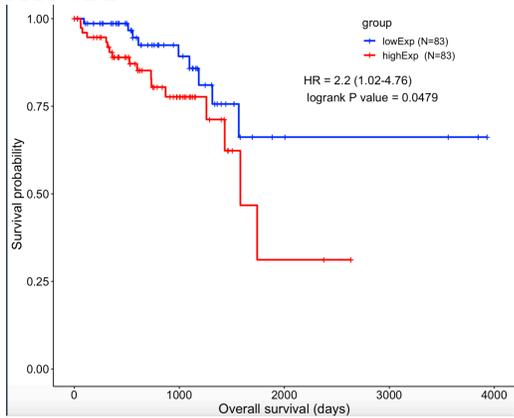


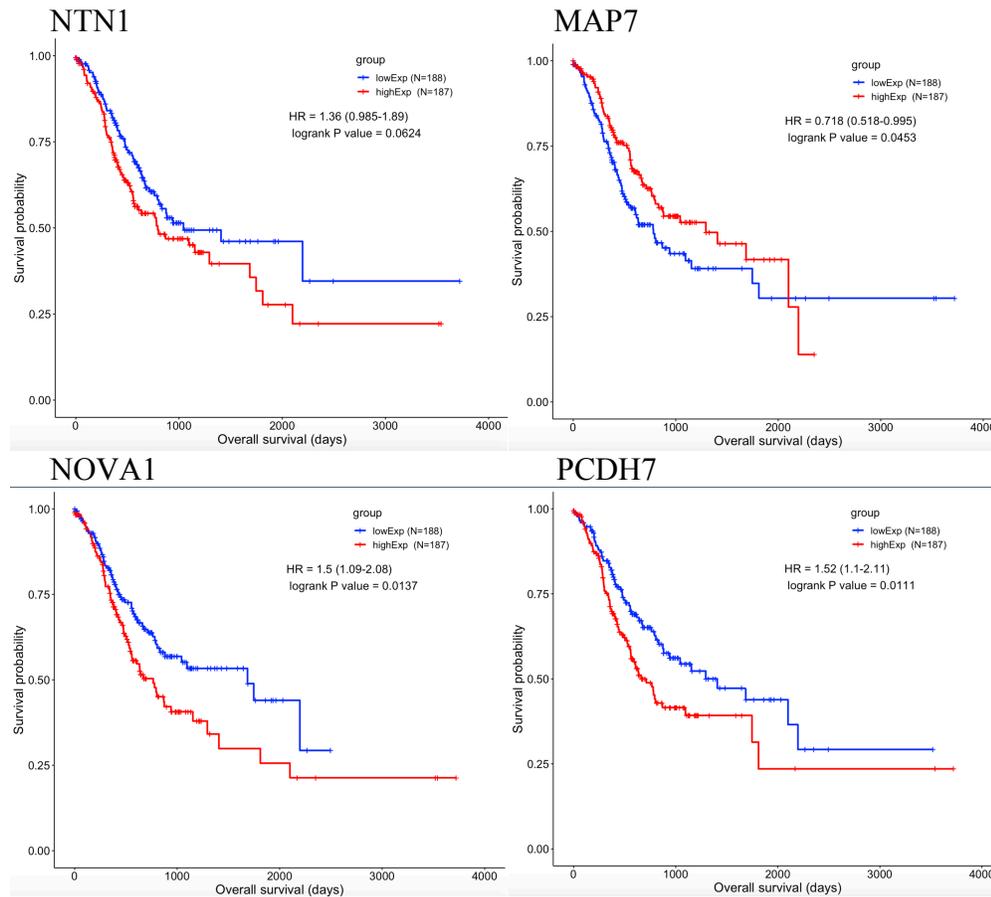
COL15A1



(C) READ

ATP8B2



(D) STAD

(E) COAD
