

Figure S1.

## Univariate cox regression in the TCGA-LAML cohort (n=151)

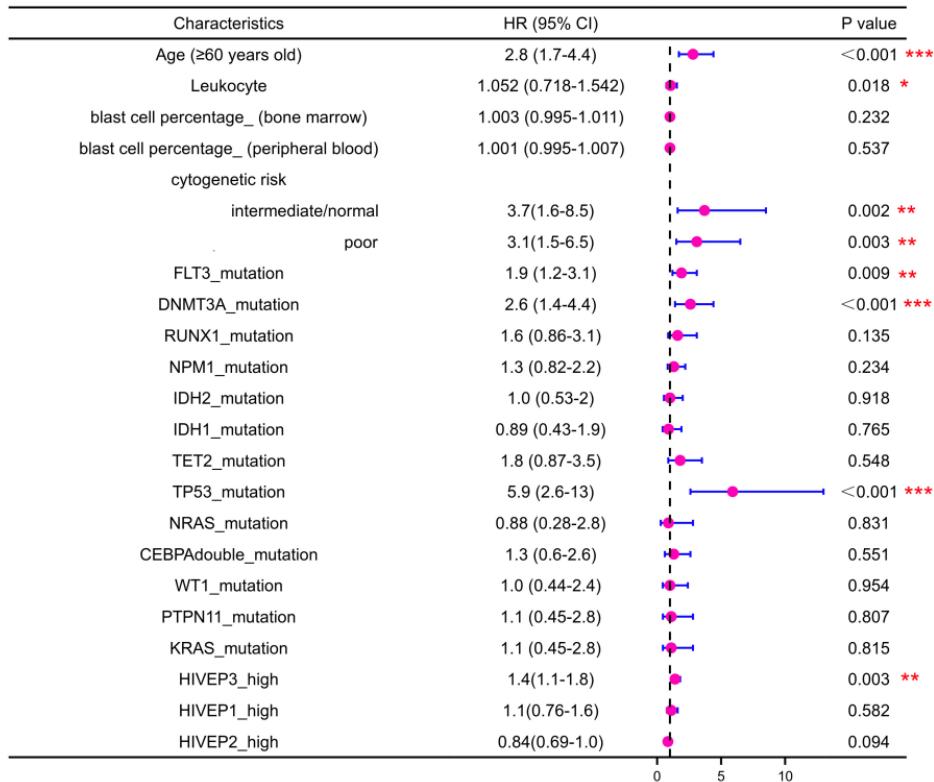


Figure S1.

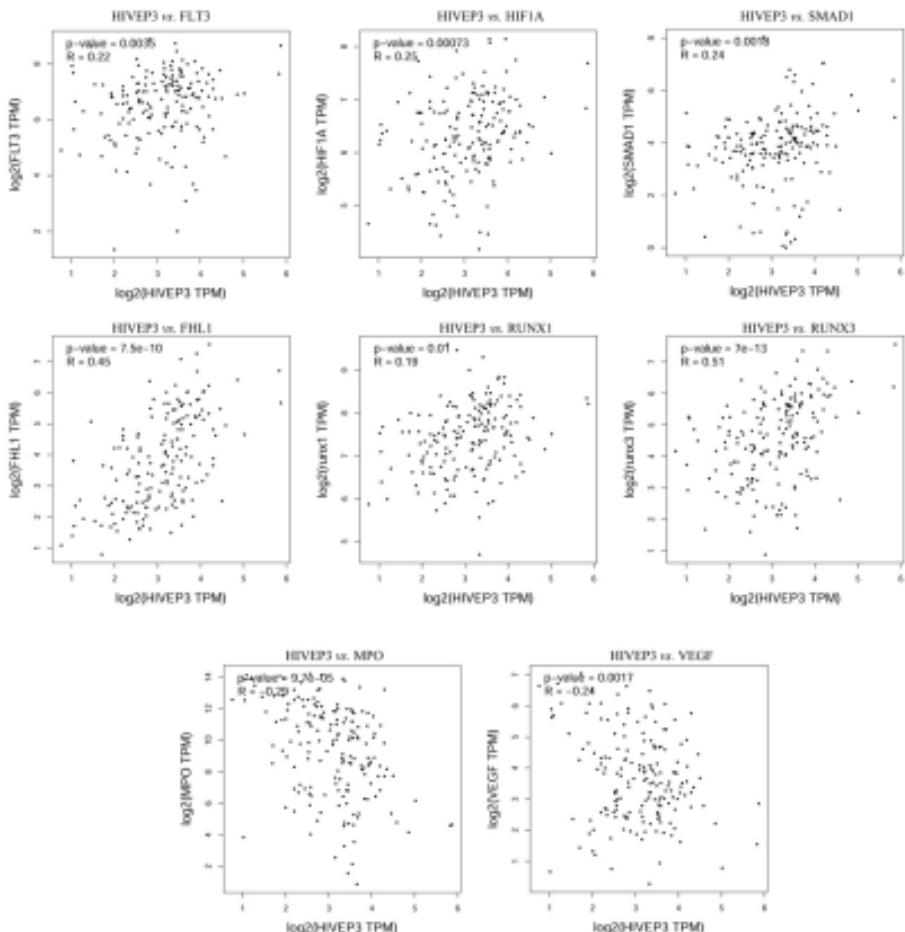
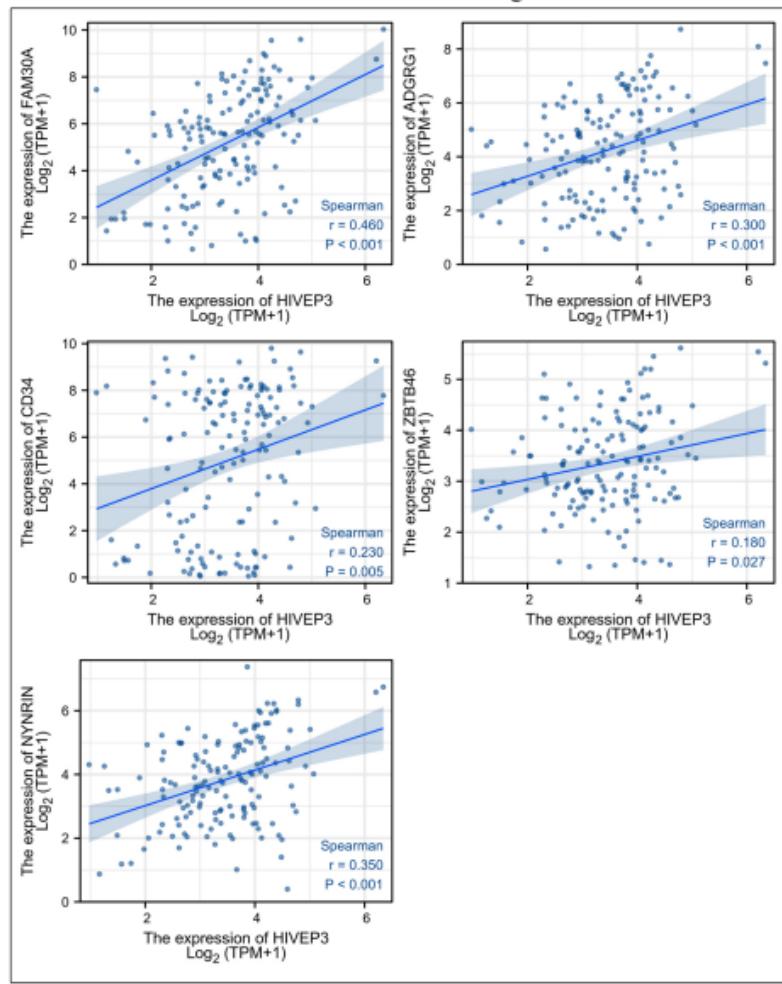


Figure S3.

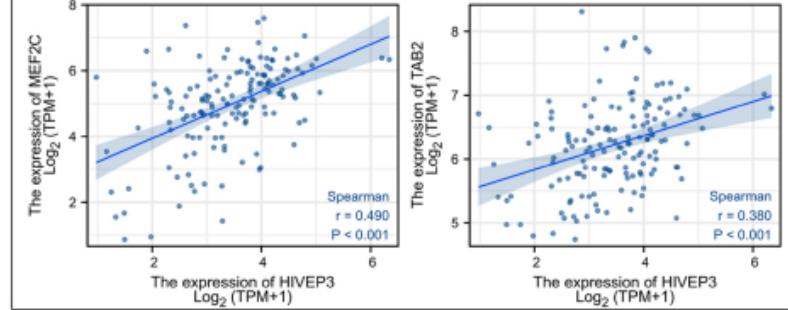
A.

HIVEP3 vs. LSCs-related genes



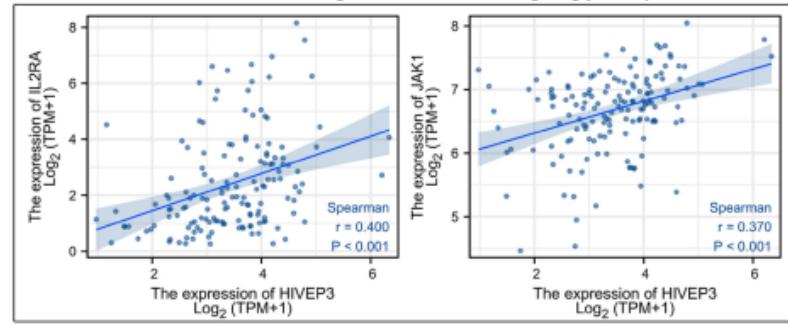
B.

HIVEP3 vs. genes in MAPK singaling pathway



C.

HIVEP3 vs. genes in JAK/STAT singaling pathway



D.

HIVEP3 vs. genes in Wnt singaling pathway

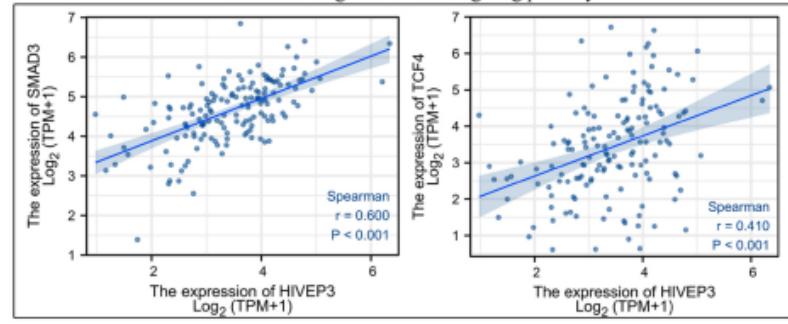
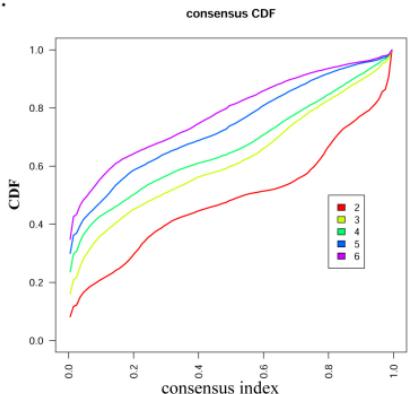
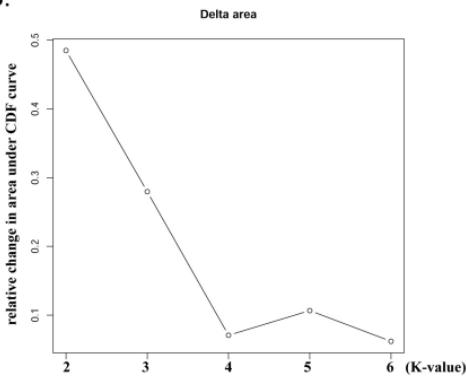


Figure S4.

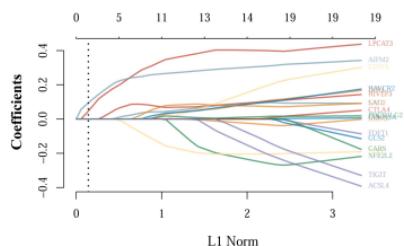
A.



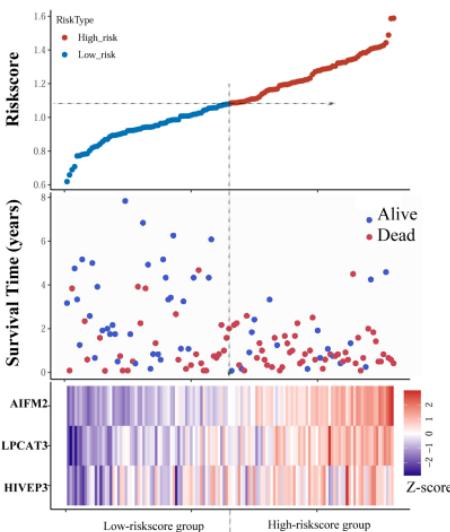
B.



C.



E.



D.

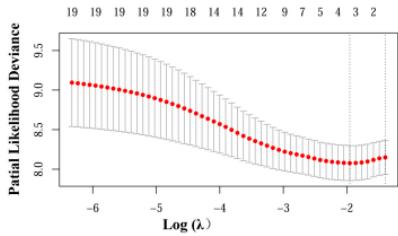
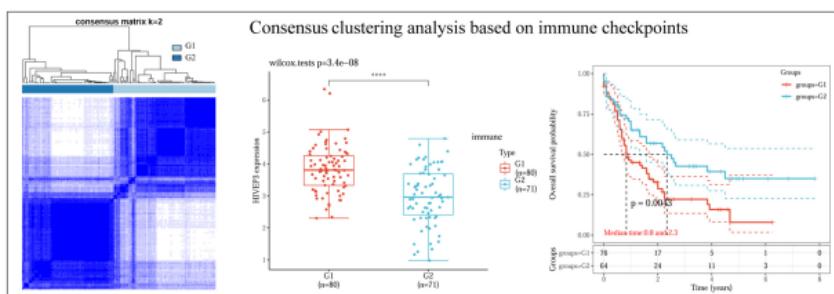
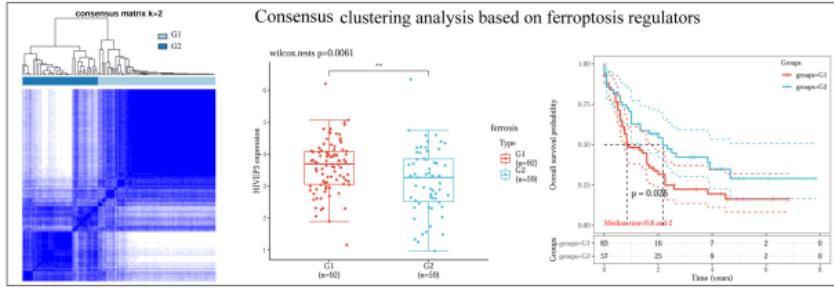


Figure S5.

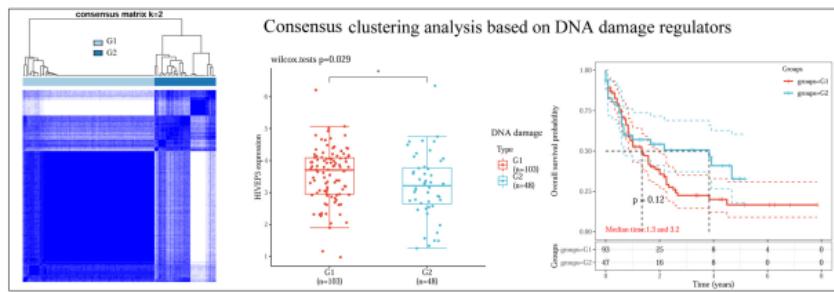
A.



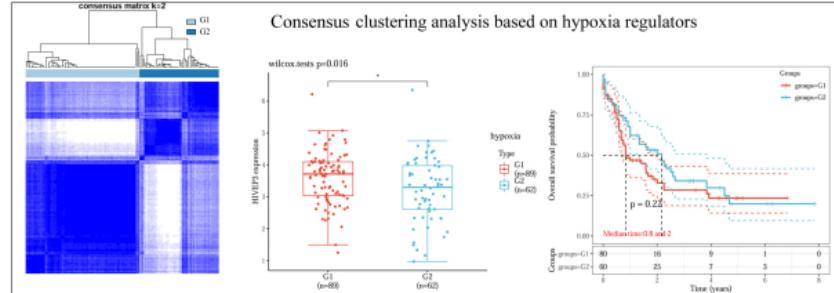
B.



C.



D.



E.

