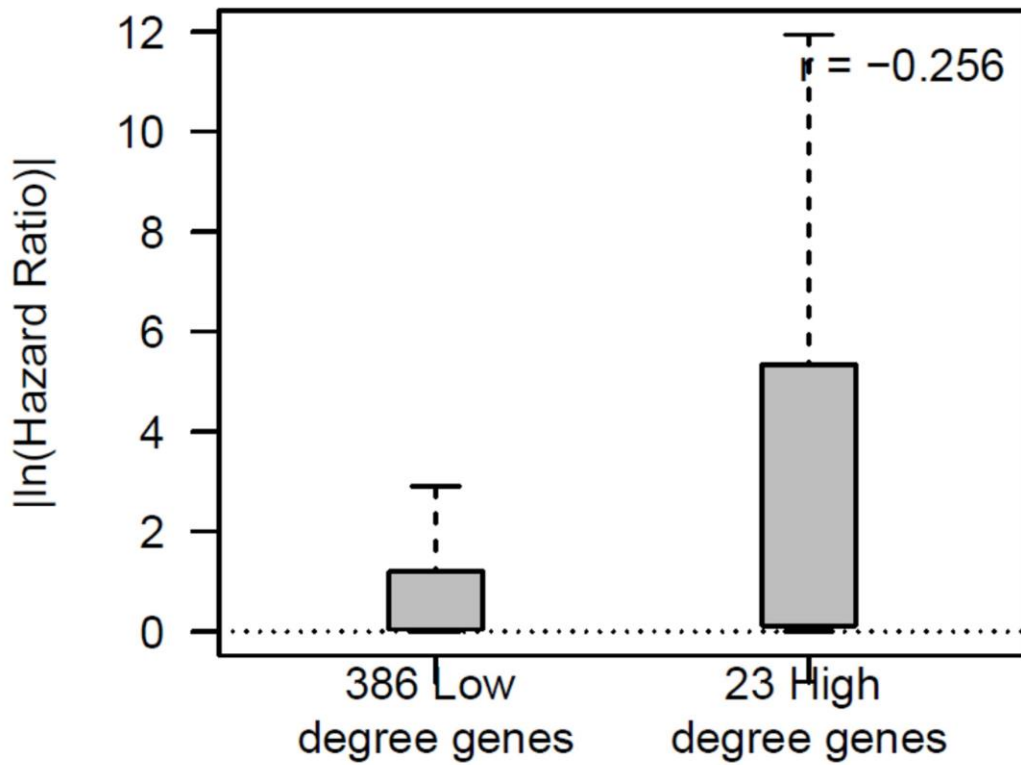


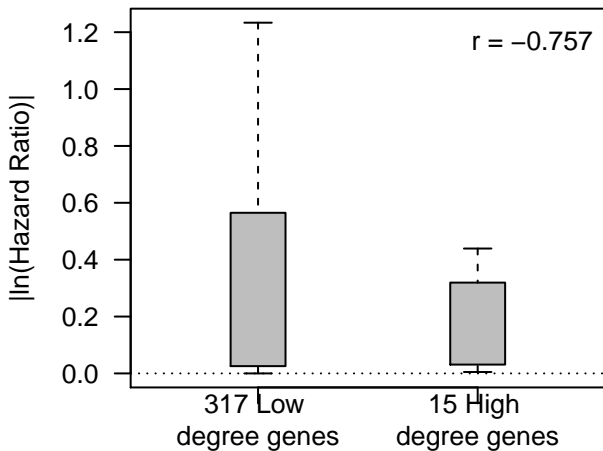
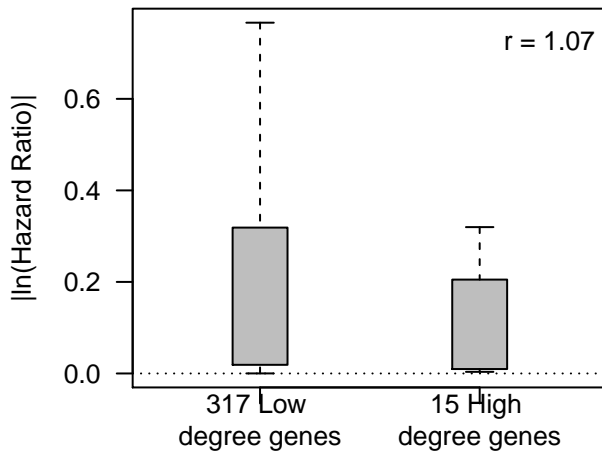
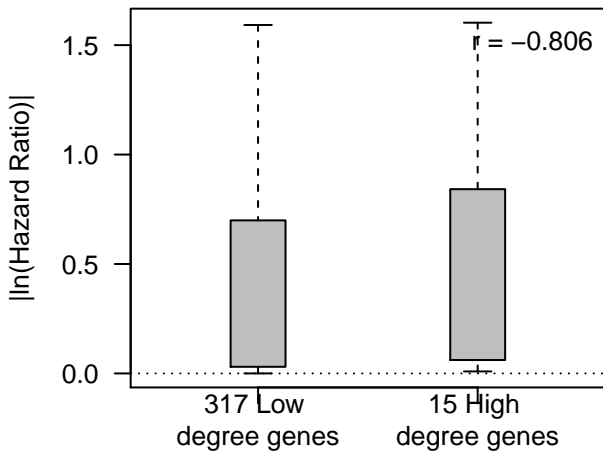
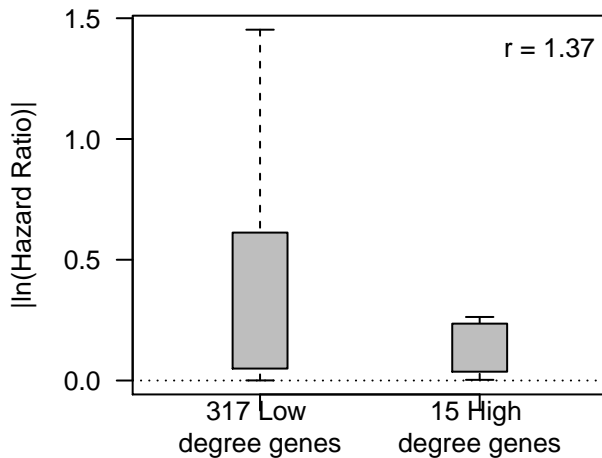
Computing microRNA-gene interaction networks in pan-cancer using miRDriver

Banabithi Bose, Matthew Moravec, and Serdar Bozdag

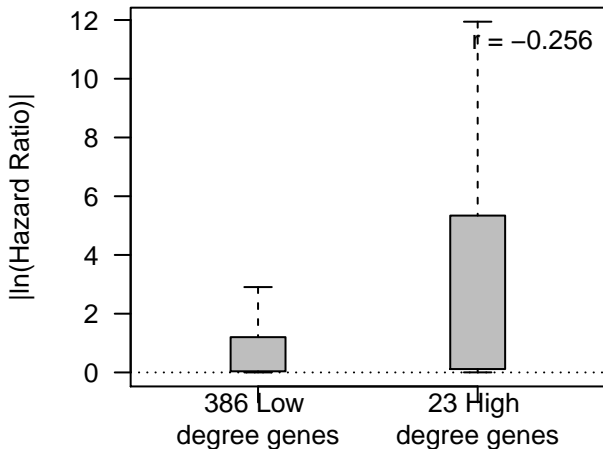
Supplemental Figure S20



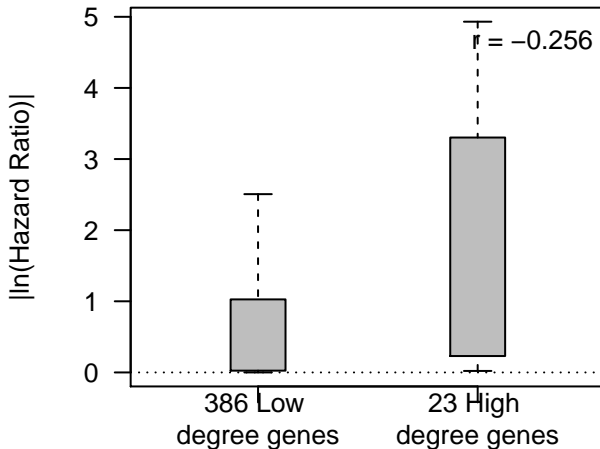
Boxplots of absolute values of natural logarithm of hazard ratios in high-degree and low-degree genes with r value of Mann-Whitney test.

BLCA OS**BLCA PFI****BLCA DSS****BLCA DFI**

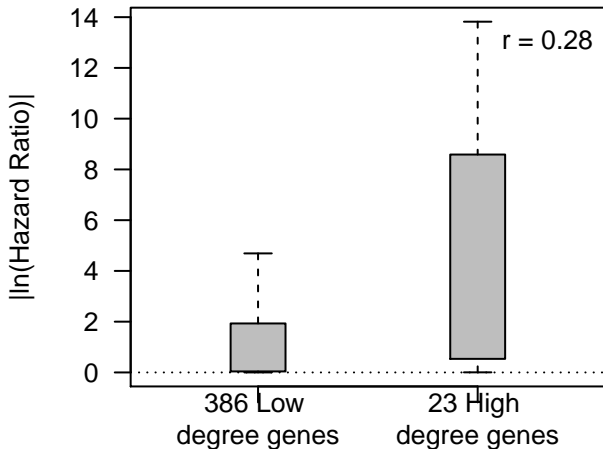
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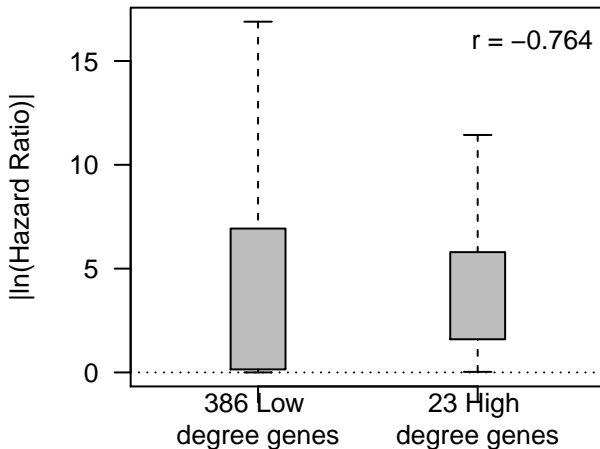
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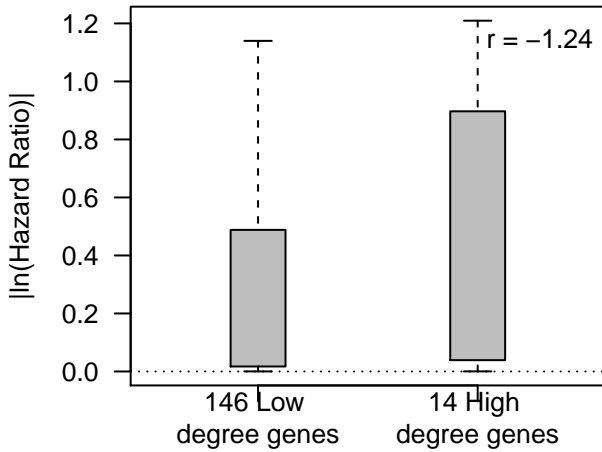
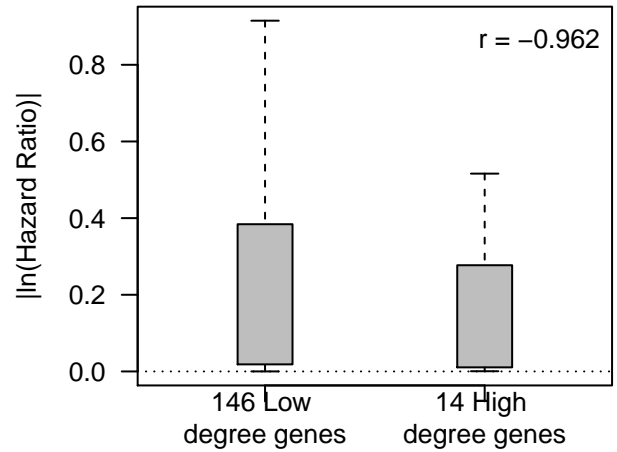
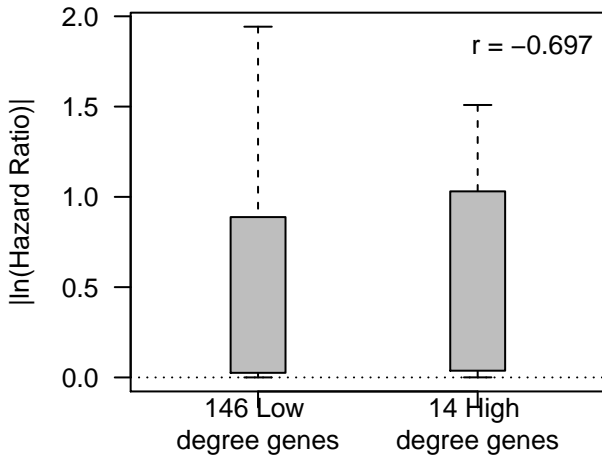
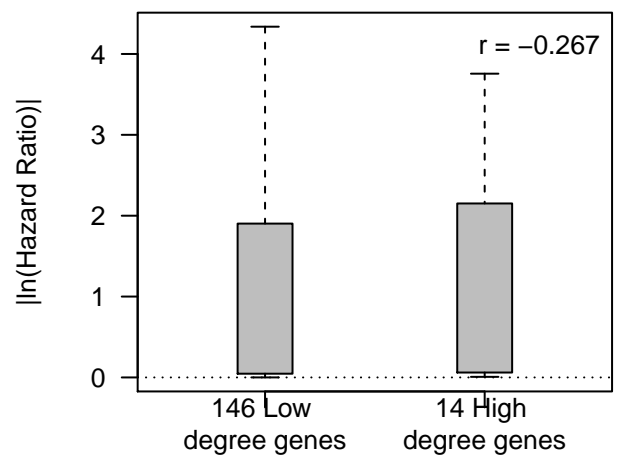


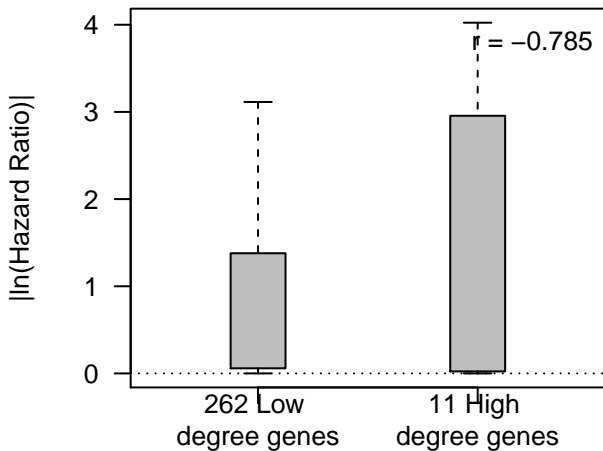
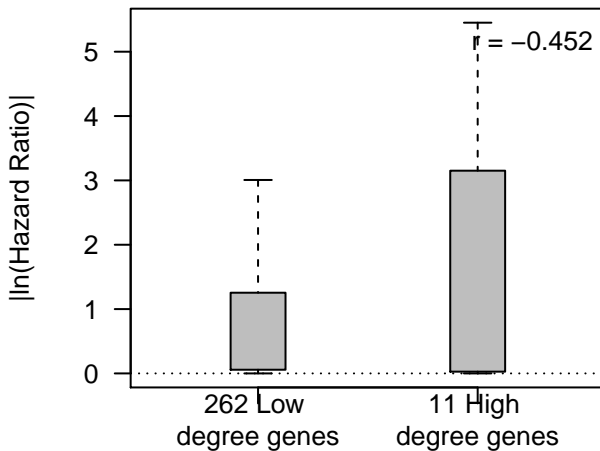
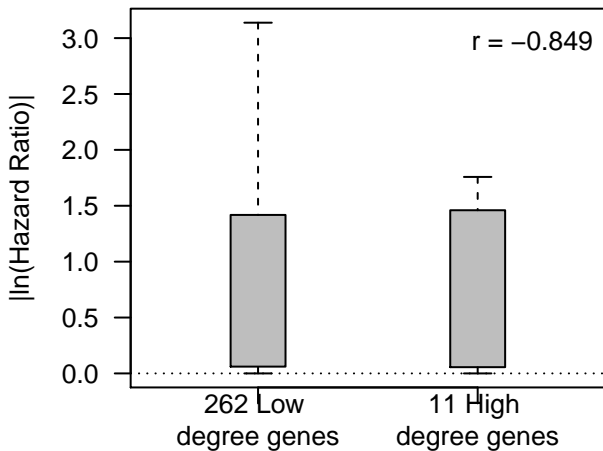
ESCA DSS



ESCA DFI



HNSC OS**HNSC PFI****HNSC DSS****HNSC DFI**

PAAD OS**PAAD PFI****PAAD DSS****PAAD DFI**