Supplementary figure

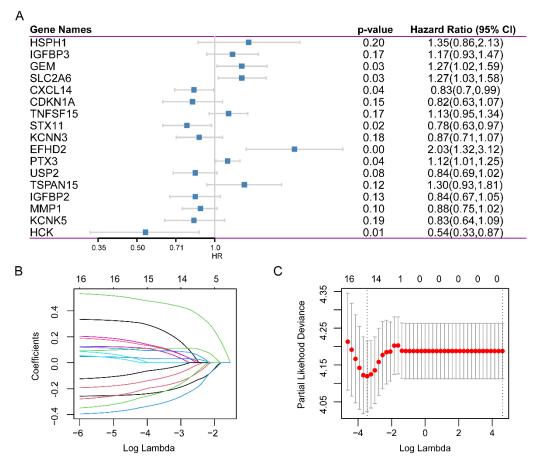


Figure S1. Identification of prognostic genes. **(A)** The univariate Cox analysis screened 17 prognostic CAFs senescence-related genes. **(B)** The LASSO regression analysis. **(C)** The partial likehood deviance of the signature.

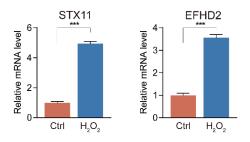


Figure S2. qRT-PCR analysis of the expression of *STX11* and *EFHD2* in control and H_2O_2 -treated MRC-5 cells. ***, p < 0.001.