

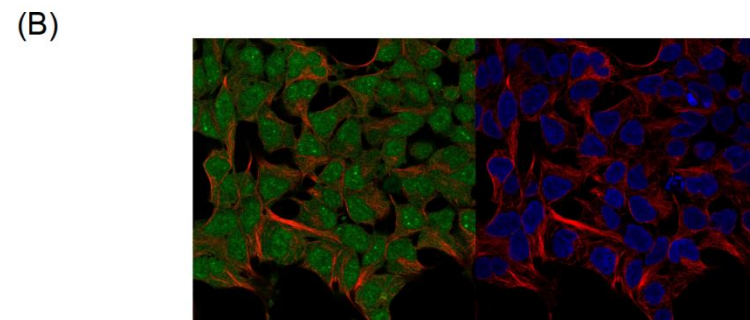
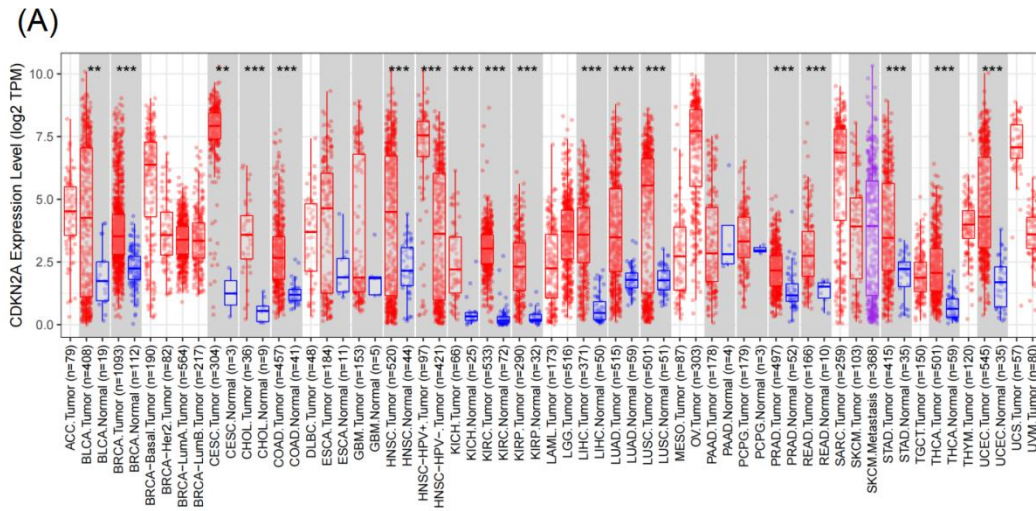
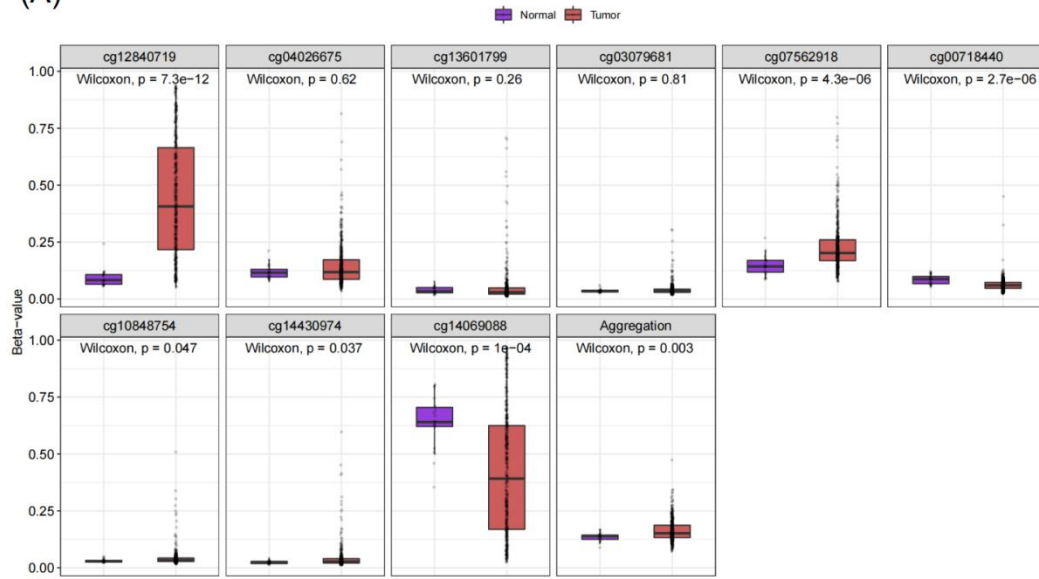
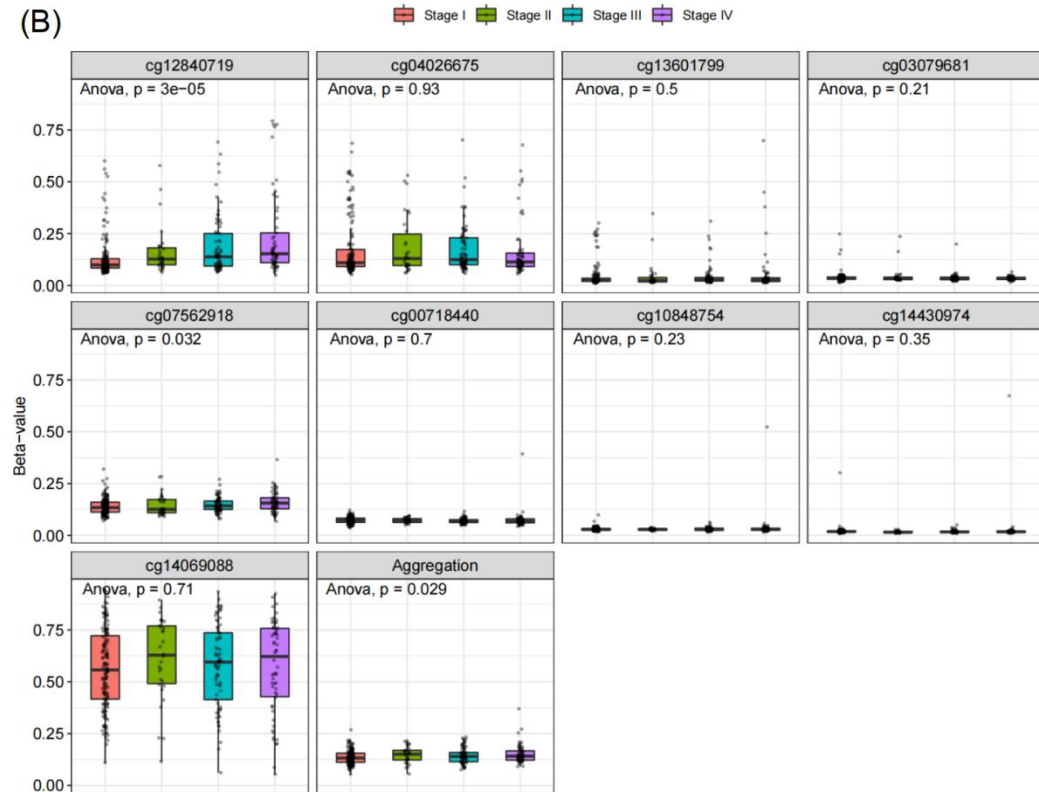


FIGURE S1 Expression and subcellular localization of CDKN2A in pan-cancer. (A) Expression of CDKN2A in pan-cancer. (B) Subcellular localization of CDKN2A. green represents the target protein, blue represents the nucleus, and red represents the microtubules.

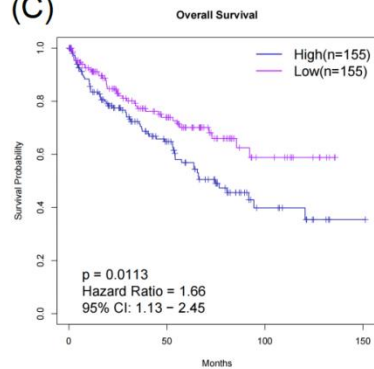
(A)



(B)



(C)



(D)

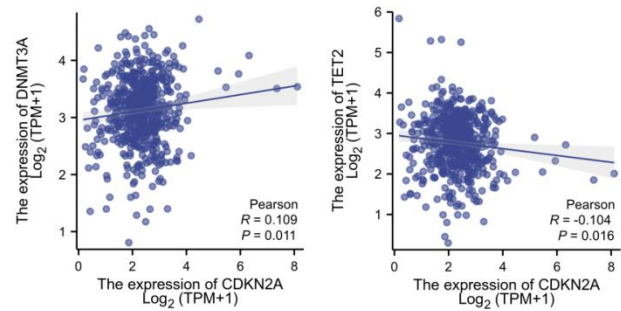


FIGURE S2 Methylation analysis of CDKN2A in ccRCC. (A) Methylation levels at 10 methylation sites of CDKN2A in ccRCC. (B) Methylation levels of 10 methylation sites of CDKN2A in ccRCC at different TNM stages. (C) Survival analysis of cg12840719. (D) Expression correlation.

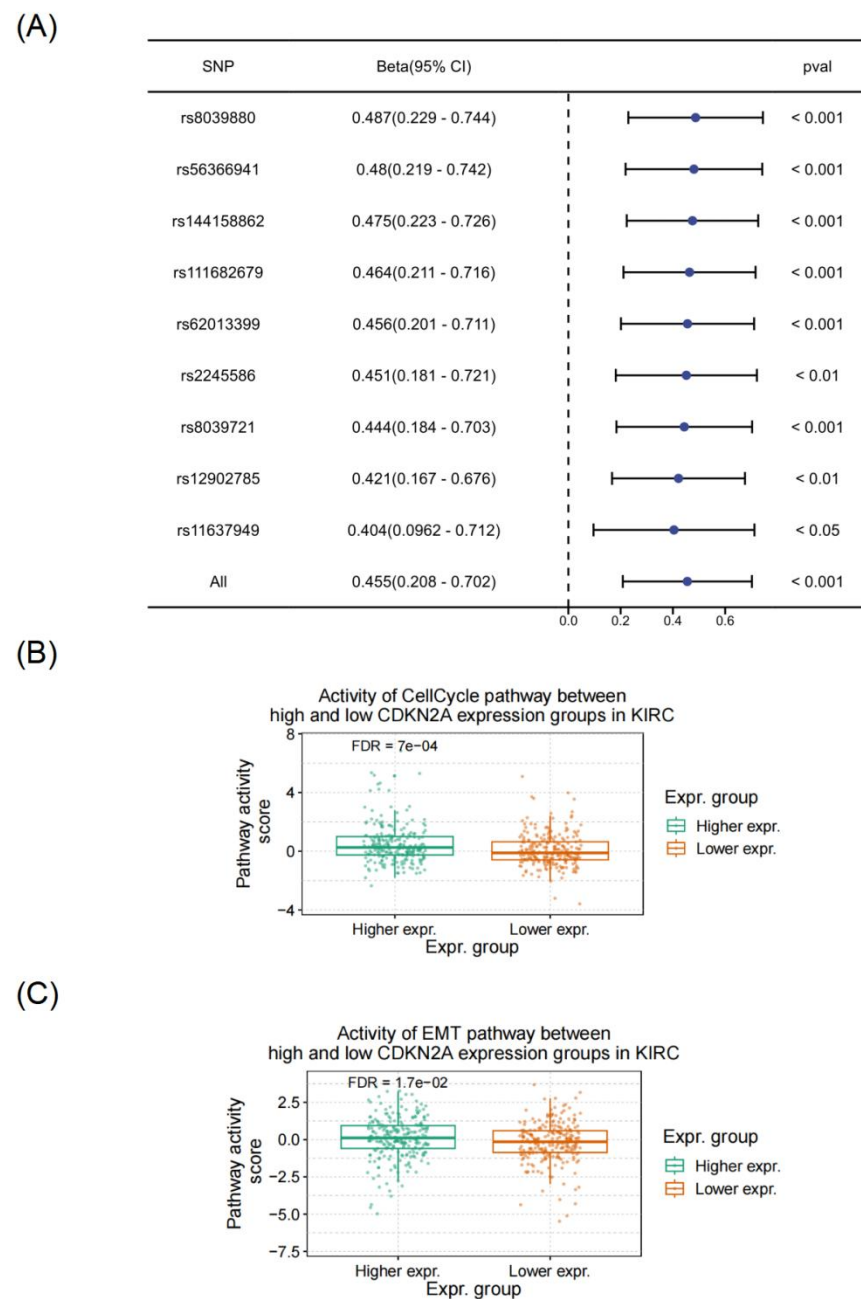


FIGURE S3 Heterogeneity test and functional exploration. (A) leave-one-out method. (B-C) Functional exploration of CDKN2A in GSCA.