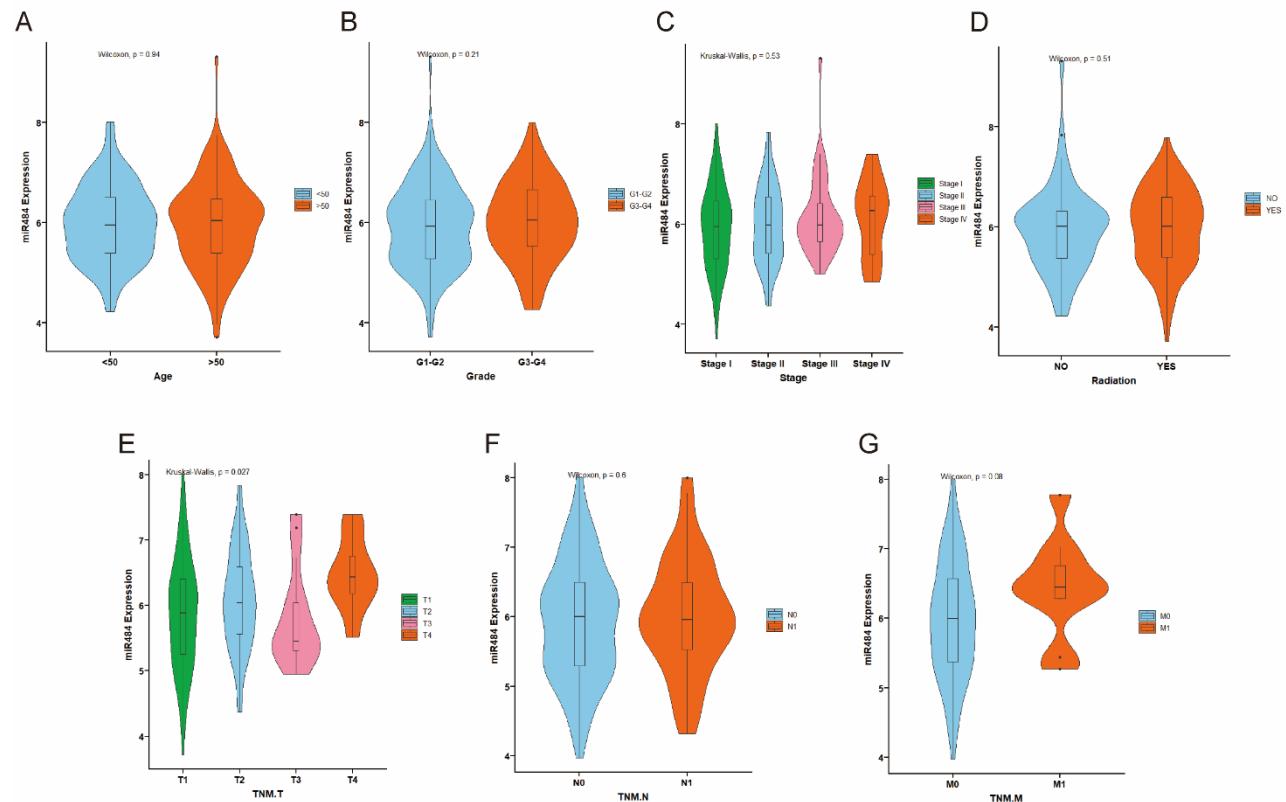
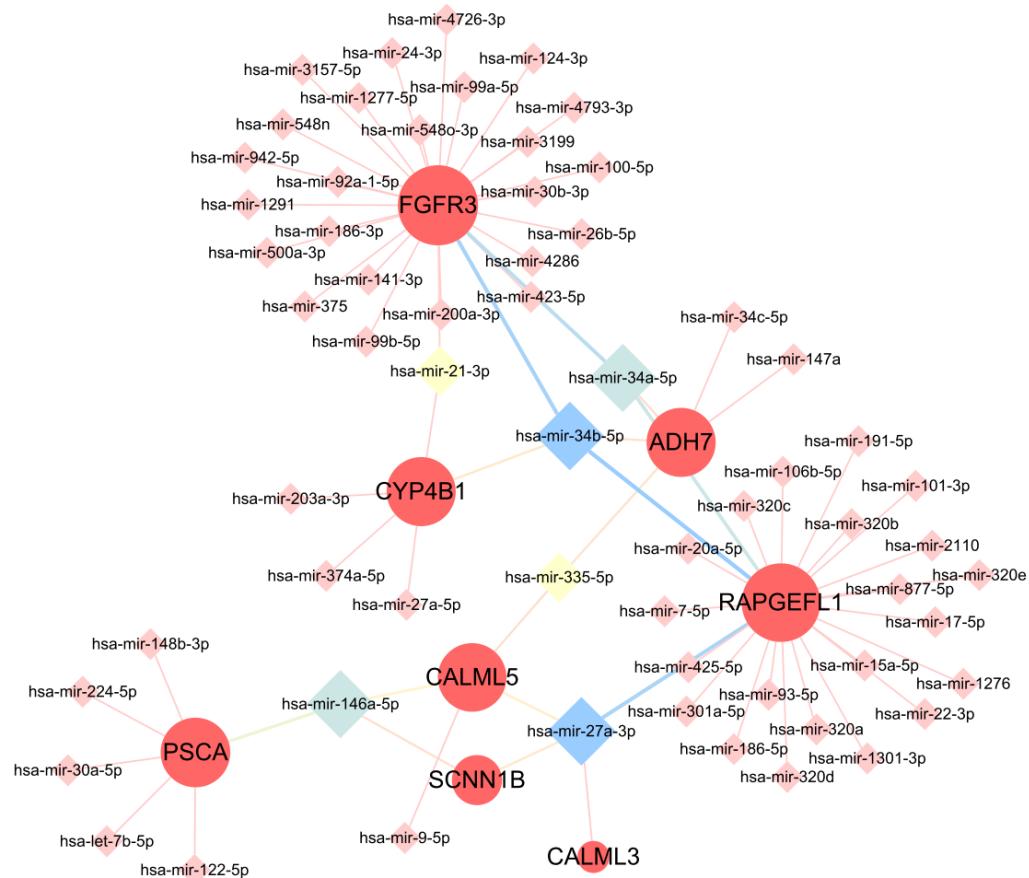


Supplementary Figure S1. Relationships between miR-484 expression and clinicopathological parameters in 283 CC tissues. (A) Age \leq 50 and age >50. (B) Grade: G1- G2 and G3-G4. (C) Stage I/I/III/ IV. (D) With and without radiation. (E-G) TNM stage.



Supplementary Figure S2. The protein-protein interaction (PPI) network presents NFs and miRNAs interacting with miR-484-related prognostic genes screened by Kaplan-Meier analysis. (A) The interaction network of prognostic genes with miRNAs. (B) The interaction network of prognostic genes with NFs.

(A)



(B)

