

SUPPLEMENTARY MATERIAL

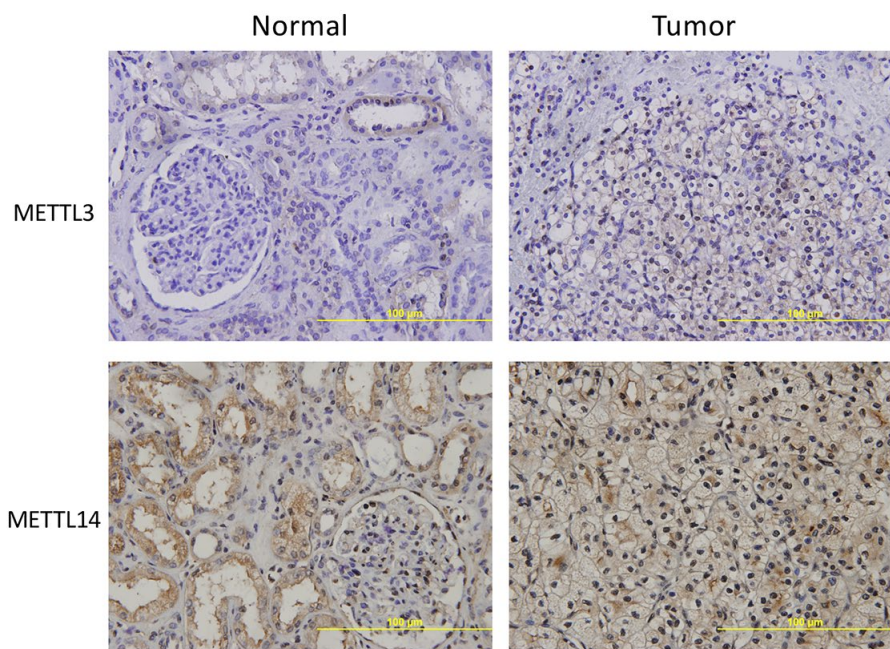


Figure S1. The protein expression of METTL3 and METTL14 in ccRCC and normal kidney tissues.

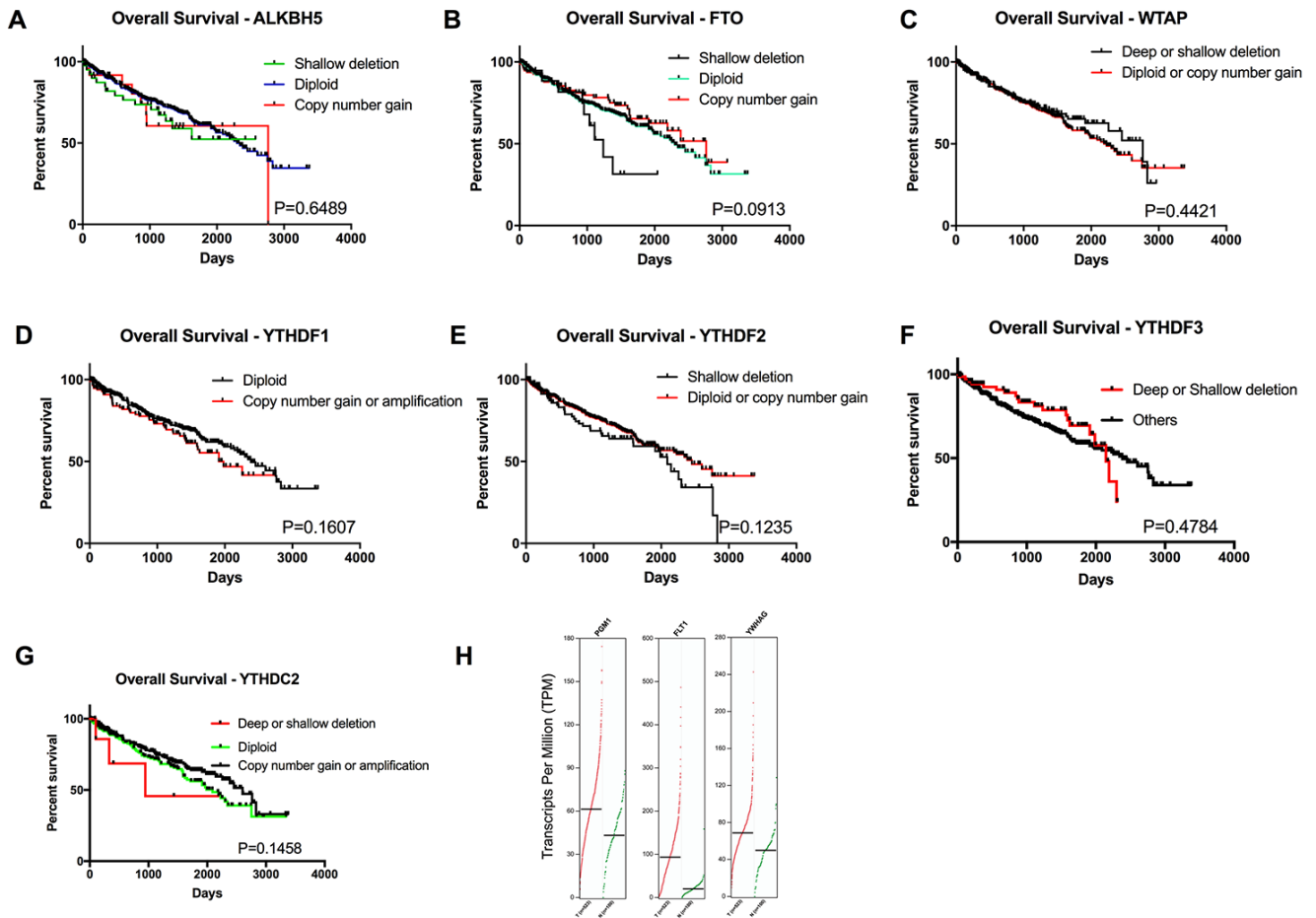


Figure S2. (A-G) OS of ccRCC patients with different alterations of m6A regulatory genes ALKBH5, FTO, WTAP, YTHDF1, YTHDF2, YTHDF3 and YTHDC2. (H) Expression of genes related to adipogenesis and mTORC1 signaling pathways in ccRCC tissues.