

Figure S1. Mutation rate of two genes in hepatocellular carcinoma using the cBioPortal database. (A and B) OncoPrint plot of MK5-AS1 and MK5 alterations.

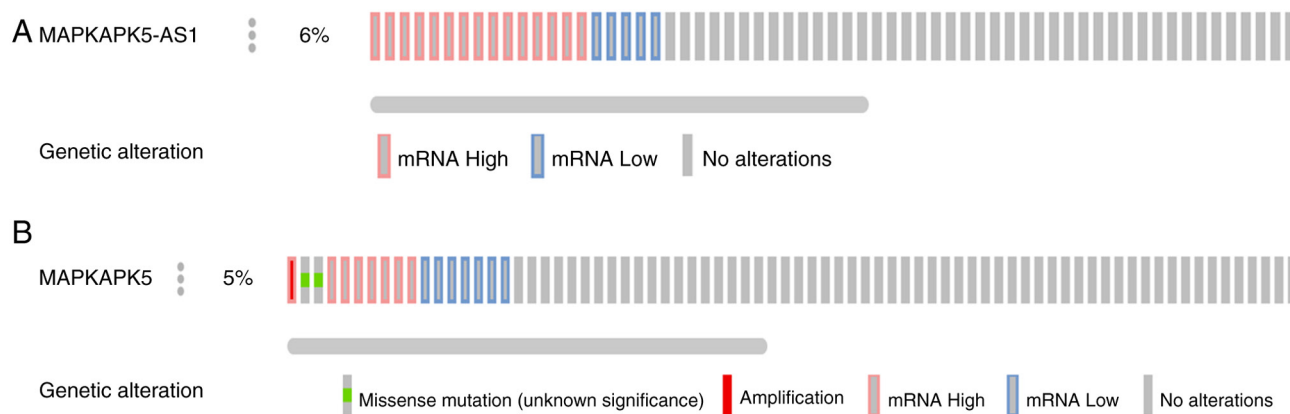


Figure S2. Expression pattern of four candidate miRs in HCC and the correlation with MK5-AS1 and MK5 using the ENCORI database. (A-D) The expression of hsa-miR-556-3p, hsa-miR-452-5p, hsa-miR-4676-3p and hsa-miR-892c-3p between HCC tissues and normal tissues. (E-L) Each picture exhibited the correlation coefficient and P-value. miR, microRNA; HCC, hepatocellular carcinoma; LIHC, liver hepatocellular carcinoma.

