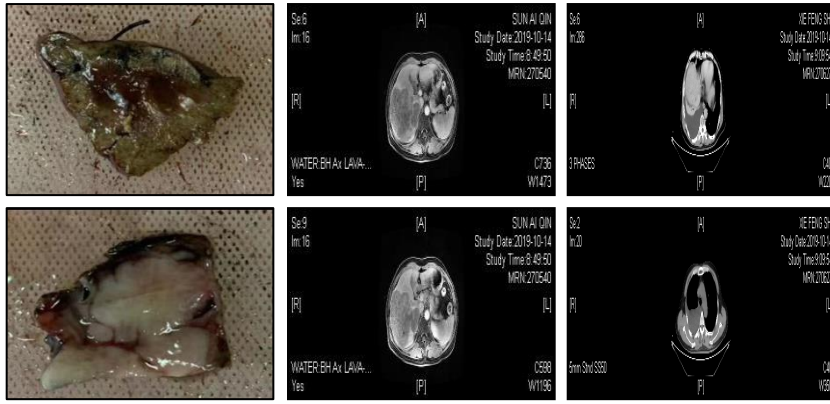


# Fig.S1

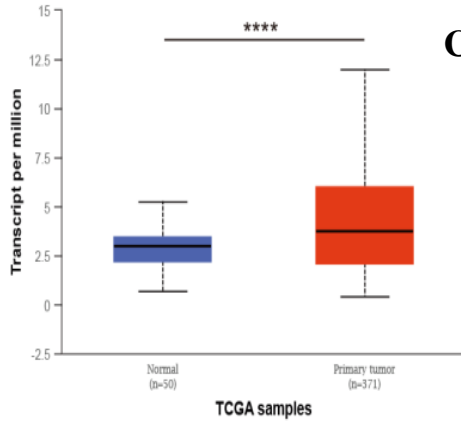
**A**

Adjacent

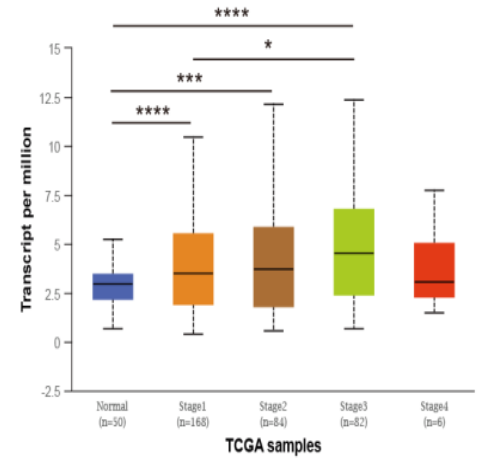
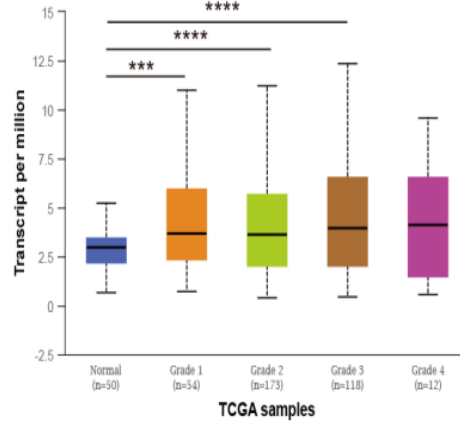


Tumor

**B**

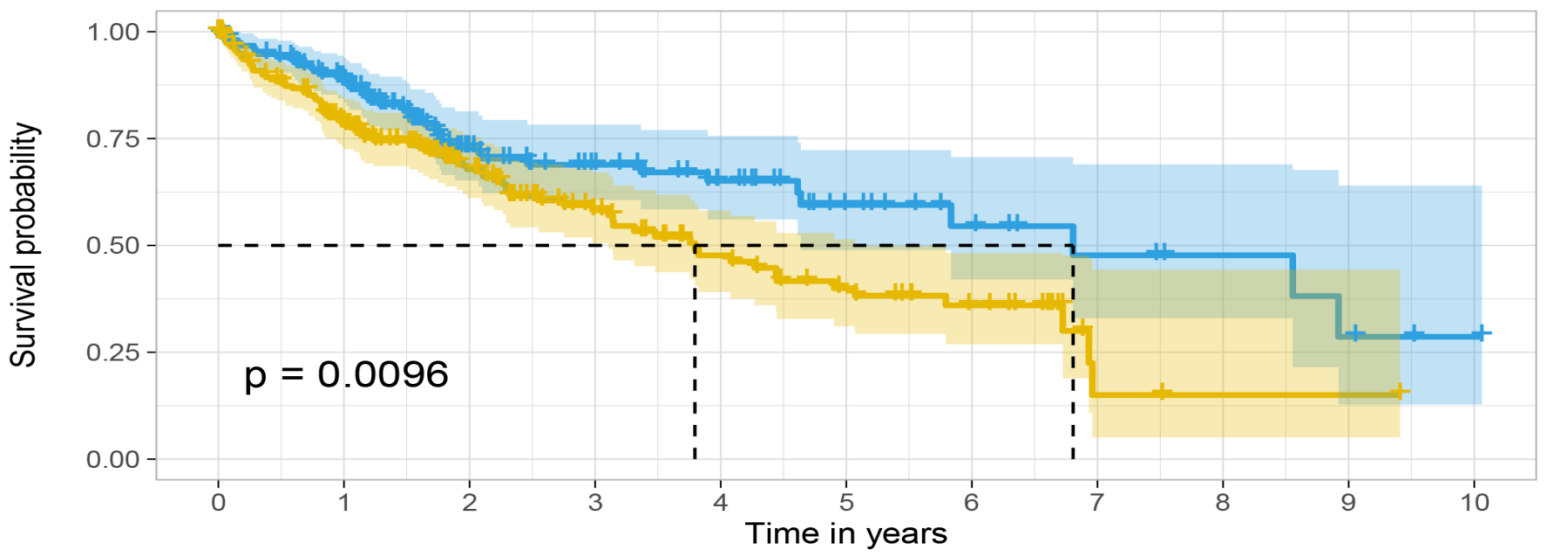


**C**

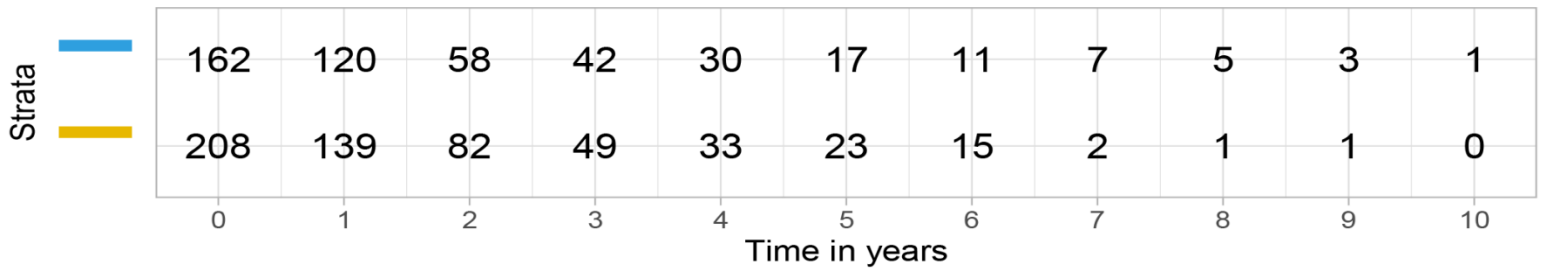


**D**

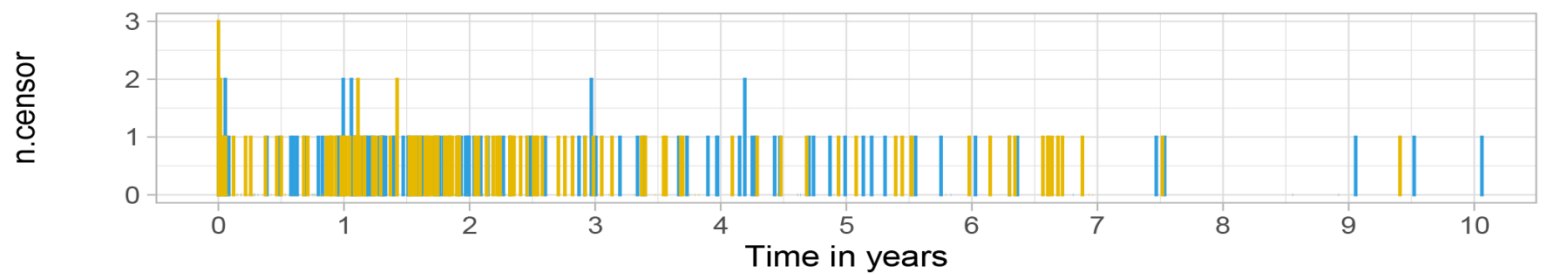
Strata ■ Low group ■ High group

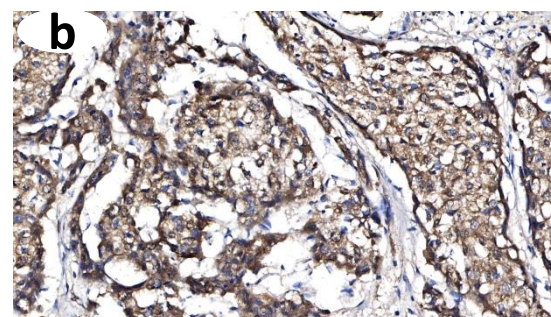
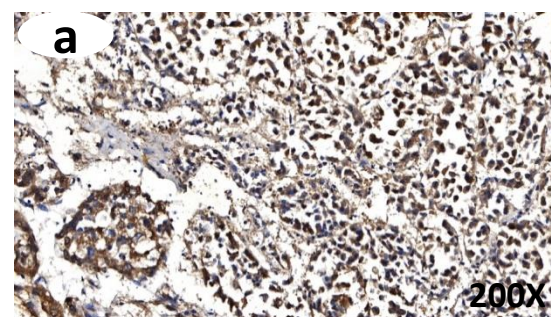
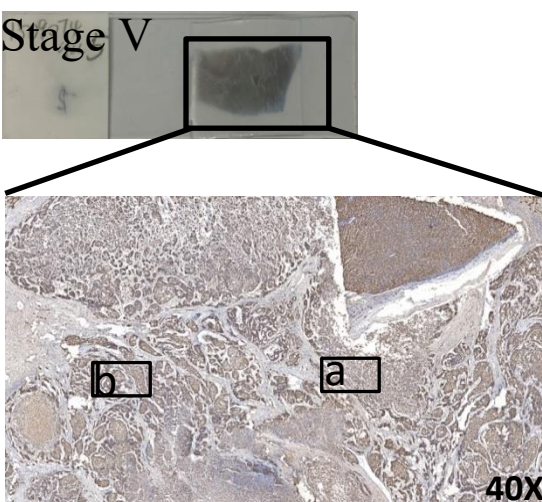
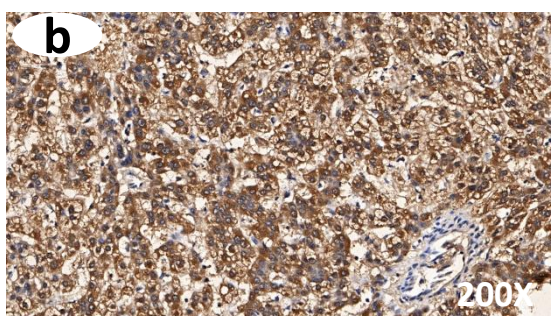
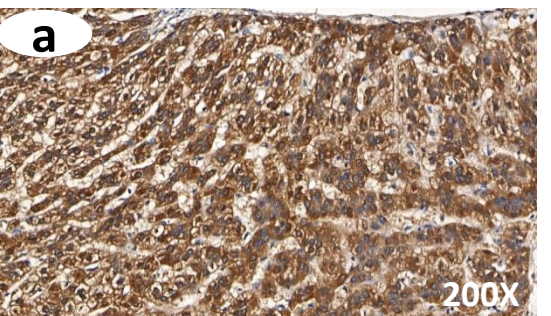
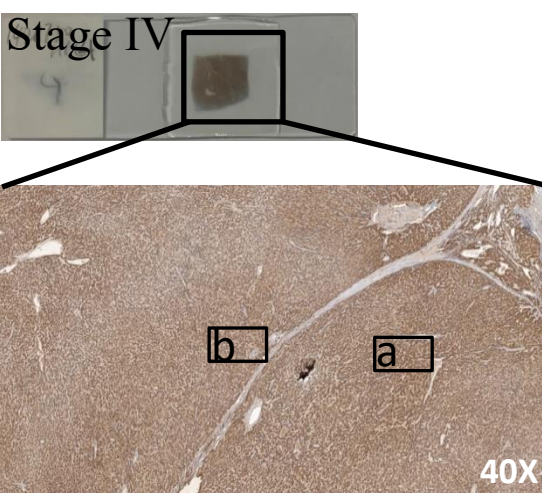
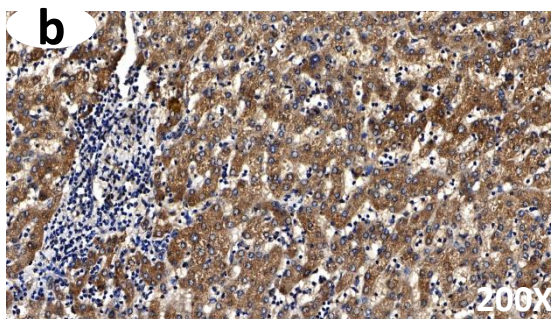
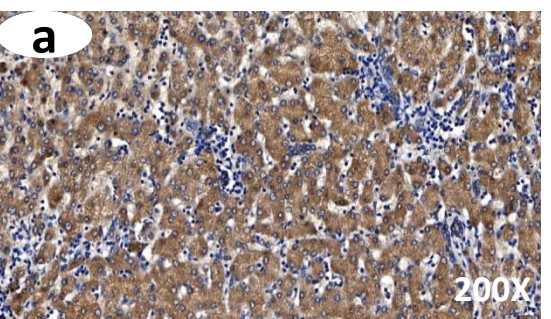
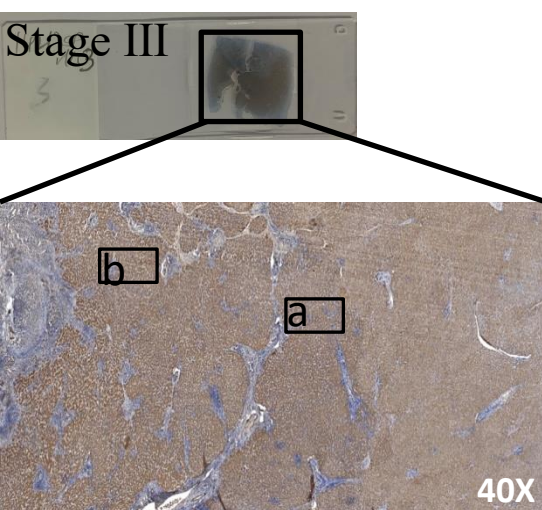
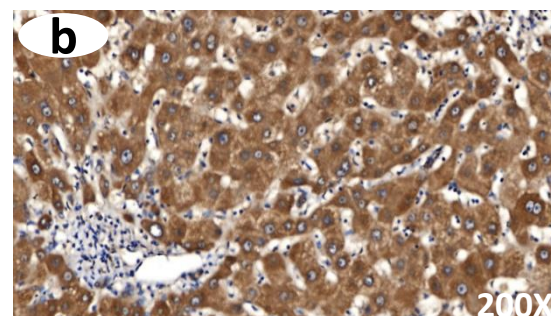
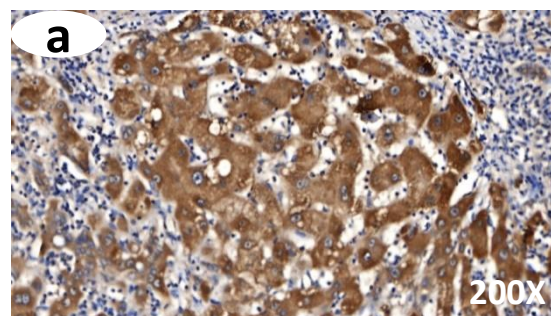
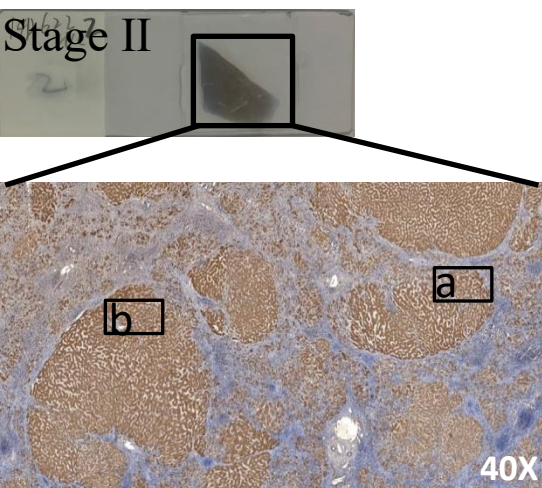
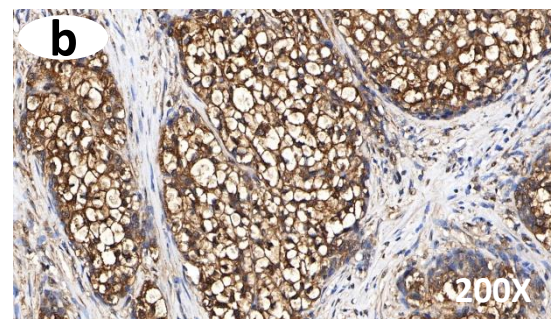
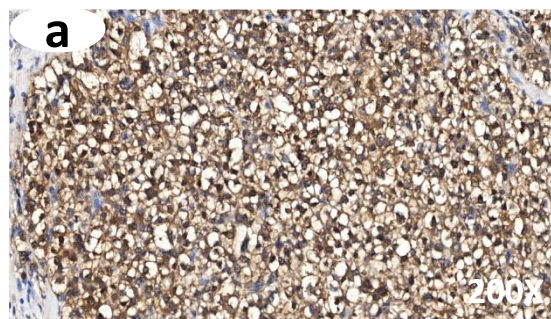
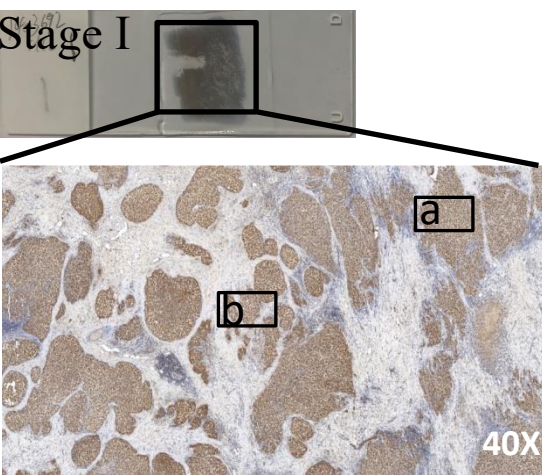


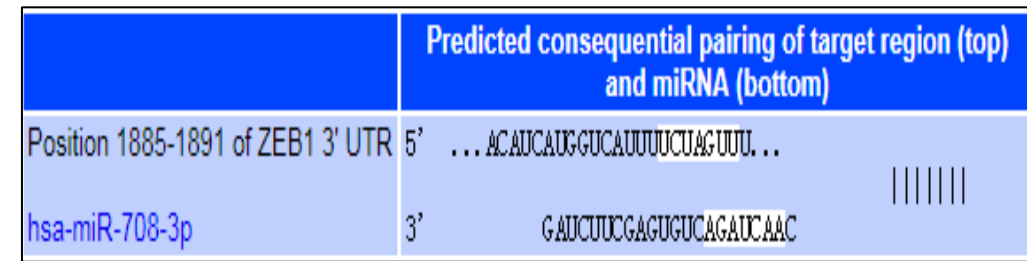
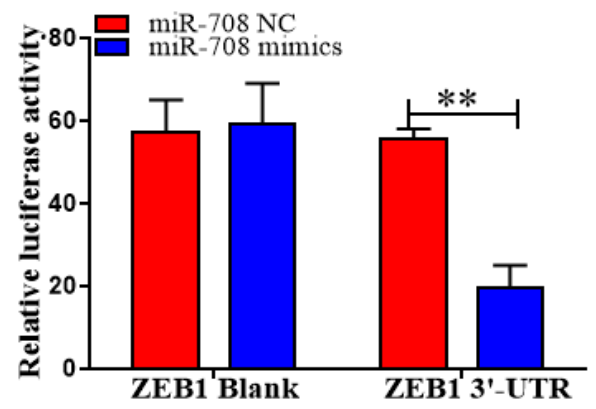
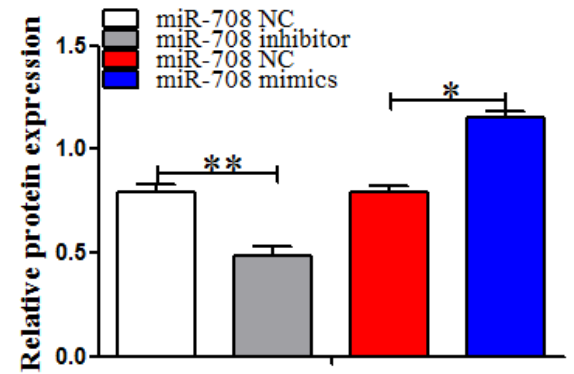
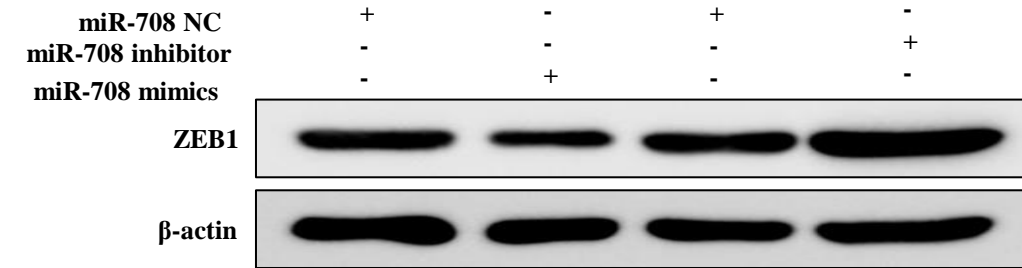
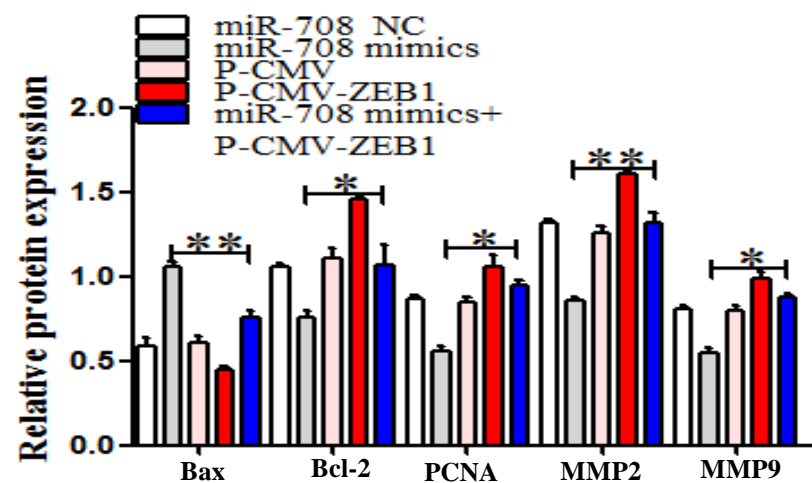
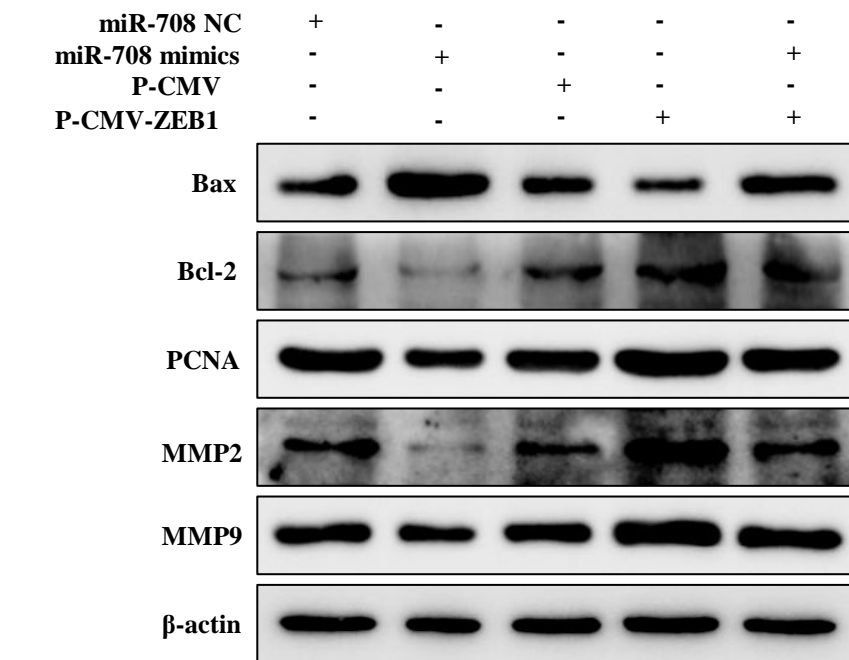
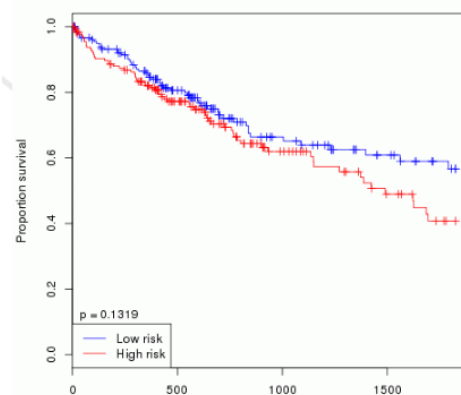
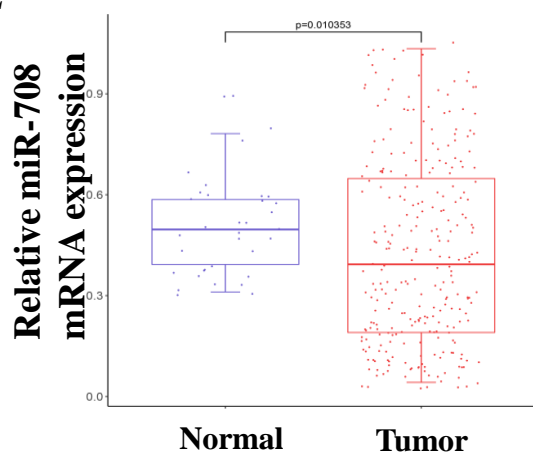
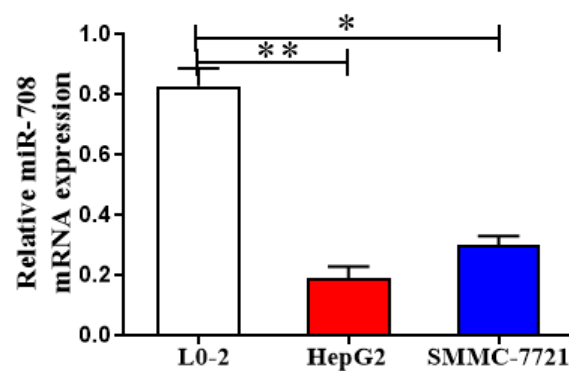
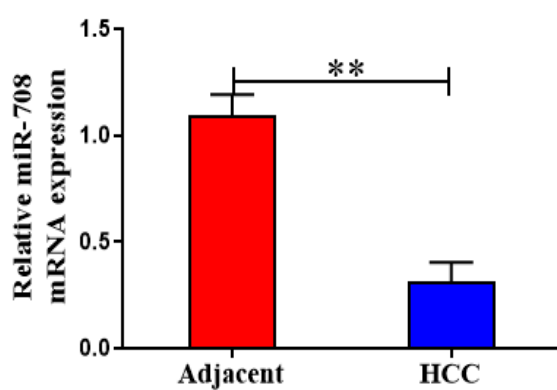
Number at risk



Number of censoring



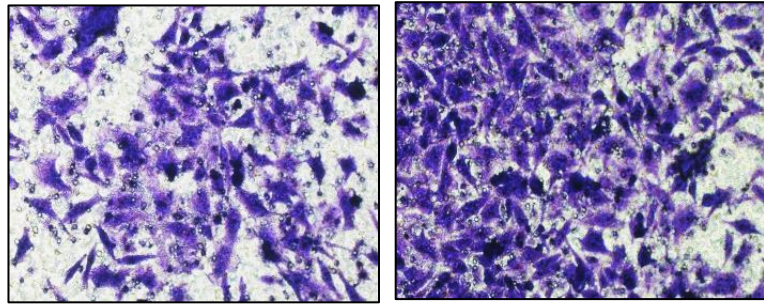
**E**

**Fig.S2****A****B****C****D****E****F**

**Fig.S3**

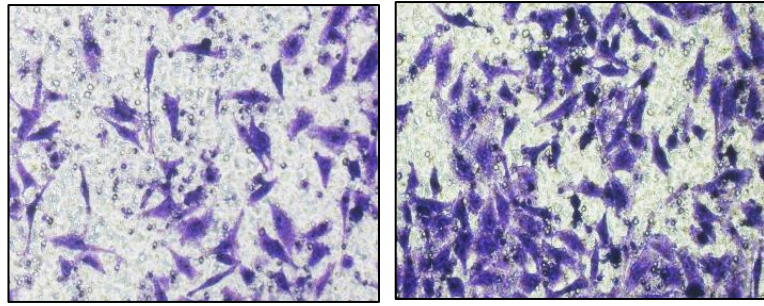
HepG2 cells

A.



miR-708 NC

miR-708 NC+TGF- $\beta$ 1



miR-708 mimics

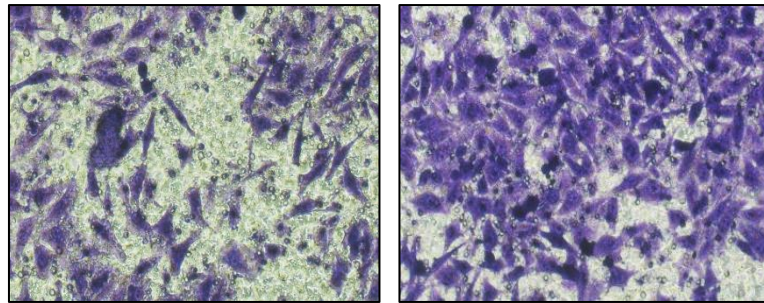
miR-708 mimics +TGF- $\beta$ 1

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**Invasion**

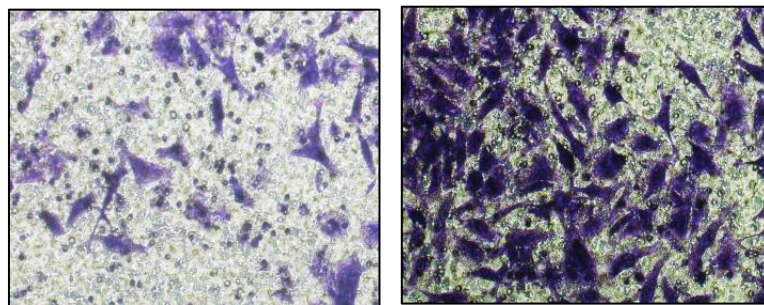
B.

HepG2 cells



miR-708 NC

miR-708 NC+TGF- $\beta$ 1



miR-708 mimics

miR-708 mimics +TGF- $\beta$ 1

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**Migration**

**Fig.S4**

