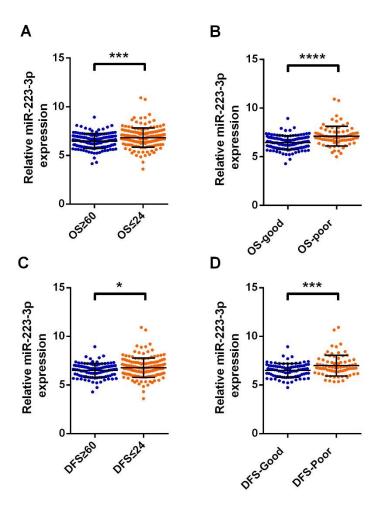


Supplementary Figure 1. External comparison of miRNAs expressions in the Cancer Genome Atlas ccRCC dataset and AUC. (A) for miR-21 and (B) for miR-452, (C) for miR-543 and (D) for miR-708. \*\* P < 0.01; \*\*\* P < 0.001, \*\*\*\* P < 0.0001.



Supplementary Figure 2. The correlation between miR-223-3p expression and long and short OS and DFS of ccRCC patients. MiRNA levels between patients with good and poor prognosis were compared by independent sample t-test. (A) for long and short OS, (B) for good and poor prognosis of OS, (C) for long and short DFS, (D) for good and poor prognosis of DFS. \* P < 0.05, \*\* P < 0.01; \*\*\* P < 0.001, \*\*\*\* P < 0.0001.