



Supplementary Figure legend. (A) The putative binding sites between miR-525-5p and circ\_0001361. (B) The predicted miR-525-5p binding sites in VMA21 3'UTR. (C) RT-qPCR was used to detect the levels of miR-525-5p and miR-520a-5p in A549 and PC9 cells. (D) The expression of VMA21 in LUAD samples and normal samples was evaluated by StarBase database. (E) The correlation between circ\_0001361 and VMA21 transcript in 32 LUAD samples was determined by Pearson correlation analysis.