



**Figure S3 Expression levels of the top eight significant miRNAs in tumor tissues and their adjacent tissues in cervical cancer patients**

(**A, B**) qRT-PCR results of miRNAs in cervical cancer tissues. Green bars indicated that miRNAs were down-regulated in sequencing data (**A**), while yellow bars indicated that miRNAs were up-regulated in sequencing data (**B**). Fold change is defined as the ratio of miRNA expression in tumor to miRNA expression in adjacent normal tissue from the same patients. Forty-six new patients were analyzed in the study.