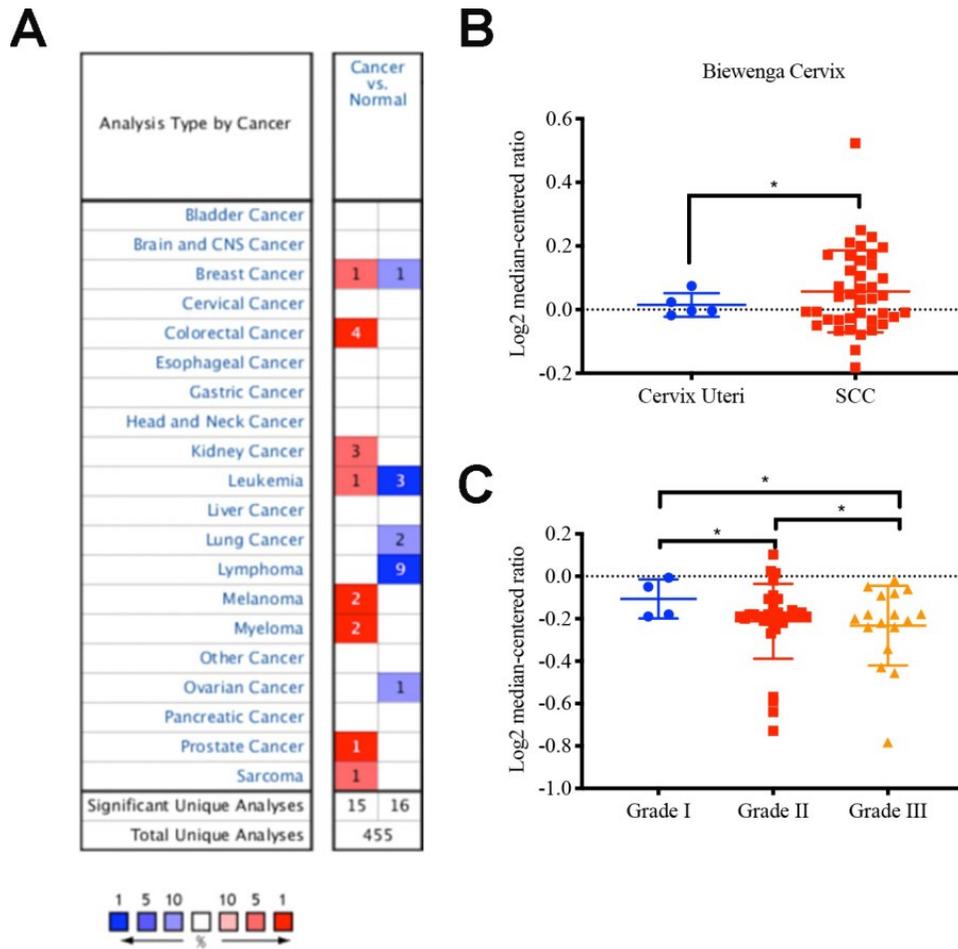
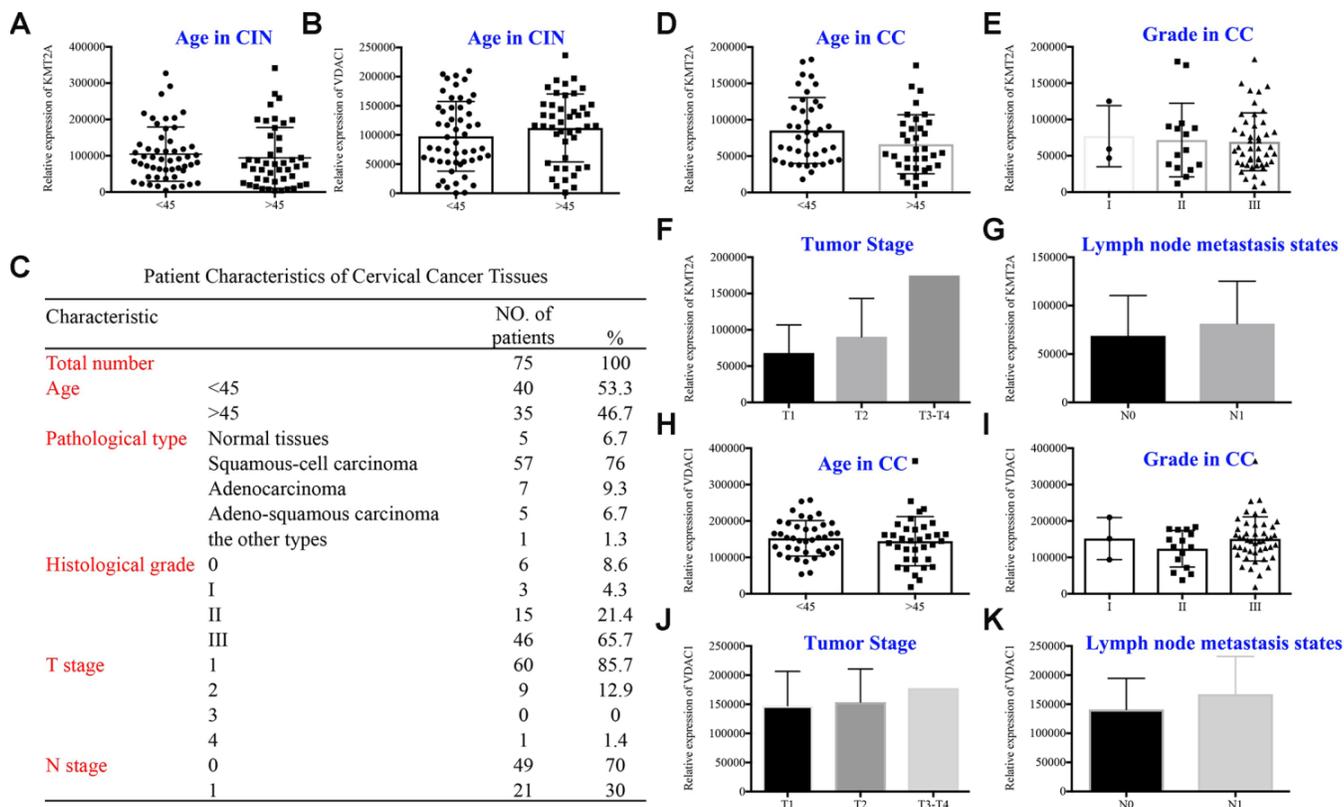


SUPPLEMENTARY FIGURES



Supplementary Figure 1. The KMT2A mRNA is upregulated in cervical cancer by analyzing Oncomine data. (A) The expression profile of KMT2A in different cancers in Oncomine data. (B) The expression of KMT2A mRNA in cervical squamous cell carcinoma and cervix uteri in Biewenga Cervix database. (C) The relationship of KMT2A mRNA and the differentiation grade of cervical cancer and cervix uteri in Biewenga Cervix database. The red color represents increase. The deeper the red is, the greater the increase. The blue represents decrease. The deeper the blue is, the greater the decrease.



Supplementary Figure 2. The clinical characteristics of cervical cancer patients. (A and B) Correlation analyses of KMT2A and VDAC1 protein expression in 102 CIN tissue samples. (D–K) Correlation of KMT2A and VDAC1 protein expression with different clinicopathologic variables in cervical cancer patient tissues.