Supplementary Figure S1. DNA methylation and gene expression of *ADCY8*, *CDH8*, *ZNF582*, and *MGMT* in the TCGA cervical cancer cohort. Correlation plot for 181 patient tumors using paired median CpG methylation (beta value) and expression value (RSEM-normalized) from each sample. Median values (Md) for each axis are marked with a dotted line. (*ADCY8*: rho = 0.01, *p* = 0.94; *ZNF582*: rho = -0.23, *p* = 0.001; *CDH8*: rho = 0.27, *p* < 0.001; *MGMT*: rho = -0.17, *p* = 0.03).