



**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*:  $\rho = -0.17$ ,  $p = 0.03$ ).