Supplementary Figure 3: Comparison of differential miRNA expression levels by p16 IHC and HPV16 ISH. The fold changes were calculated as differential miRNA expression levels between p16+ve vs. p16-ve by IHC, and HPV+ve vs. HPV-ve by ISH. The x and y axis refer to fold changes defined HPV16 ISH and p16 IHC, respectively. These two sets of p-values are highly correlated, with $R^2=0.78$. 