



Supplementary Figure 2. Correlation plots of independently significant genes identified by multivariate analysis. (A) Expression *COX16* and *COX17* were significantly correlated ($r=0.62$, $p=5.2 \times 10^{-9}$) in HPV+ HNSCCs. (B) Expression *SLC16A2* and *COX16* were not correlated ($r=-0.084$, $p=0.48$). (C) Expression of *SLC16A2* and *COX17* were not correlated ($r=0.0013$, $p=0.99$).