

Supplementary Table III. The log fold change of the genes in HPV +ve in comparison to HPV -ve head-and-neck squamous cell carcinoma (HNSCC) samples from different datasets as data mined from HNCDB database (<http://hncdb.cancerbio.info>)

Genes	TCGA (fold change)	<i>P</i>	GSE55546 (fold change)	<i>P</i>	GSE55544 (fold change)	<i>P</i>	GSE55542 (fold change)	<i>P</i>	GSE9366 (fold change)	<i>P</i>
HIF1 α	0.795	2.07E-05	0.11	7.63E-01	-0.123	1.76E-04	-0.126	3.83E-06	-0.7	7.48E-02
VHL	0.479	2.62E-03	0.033	3.40E-01	0.055	2.15E-02	0.097	1.06E-04	-0.61	3.49E-01
LIMD1	-0.068	6.57E-01	-0.064	5.72E-02	0.014	6.12E-01	-0.005	8.34E-01	1.55	2.58E-02
VEGF	0.325	1.43E-01	0.3	2.35E-04	0.118	9.00E-02	0.09	1.01E-01	-0.194	6.75E-01