

Table S10. DNA methylation β -values in different patient groups in the TCGA-HNSCC cohort (RNA-Seq). These groups were compared by t-test

		mean values				t-test p-values			
gene name	probe name	Normal	HPV+ HNSCC	HPV- HNSCC	HNSCC samples	Normal vs HNSCC	Normal vs HPV- HNSCC	Normal vs HPV+ HNSCC	HPV+ vs HPV- HNSCC
ZNF14	cg17476421	0.022708	0.165640595	0.116051	0.12245	2.61E-16	2.61E-16	0.000267	0.18017
ZNF160	cg08228914	0.025282	0.144364065	0.151199	0.150317	3.88E-18	3.88E-18	0.002679	0.862113
ZNF420	cg14340610	0.061692	0.110454937	0.221476	0.207151	3.5E-21	3.5E-21	0.027134	1.64E-05

significant p-values are bolded