



Figure S1: Gene expression reanalysis in the oropharynx. RSEM normalized RNA-seq data for tumors occurring in the oropharynx was extracted from the TCGA database for the HNSC cohort for HPV+ and HPV- samples. Statistical analysis was performed using a two-tailed non-parametric Mann-Whitney U test. Numbers in brackets refer to the number of samples included in each analysis. * $p \leq 0.05$, ** $p \leq 0.01$, *** $p \leq 0.001$, **** $p \leq 0.0001$, ns - not significant