



Supplementary Figure 5. Expression of G2/M kinetochore components in TCGA HNSCC. A) Box plot analysis of *PLK1*, *AURKA*, *TPR*, *NUF2* and *NDC80* mRNA expression in normal ($n = 44$) and HNSCC ($n = 516$) tissue from TCGA HNSCC database. **B)** Box plot analysis of *PLK1*, *AURKA*, *TPR*, *NUF2* and *NDC80* mRNA expression in normal ($n = 44$), HPV- HNSCCs ($n = 434$) and HPV+ HNSCCs ($n = 80$) tissue from TCGA HNSCC database.