

Figure S1. The prognostic value of hub genes in the overall survival of HNSCC patients based on TCGA data.

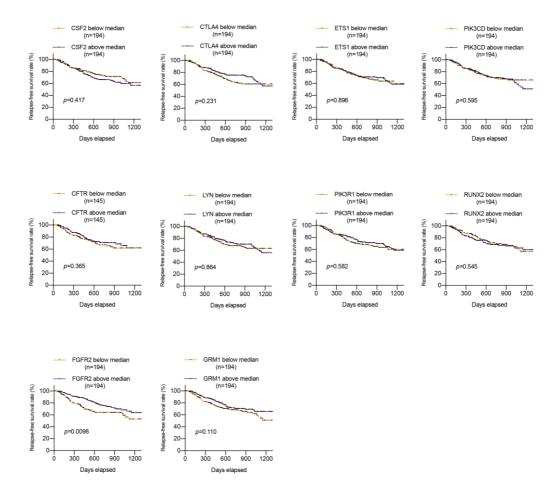


Figure S2. The relationship of hub genes with the relapse-free survival of HNSCC patients based on TCGA data.

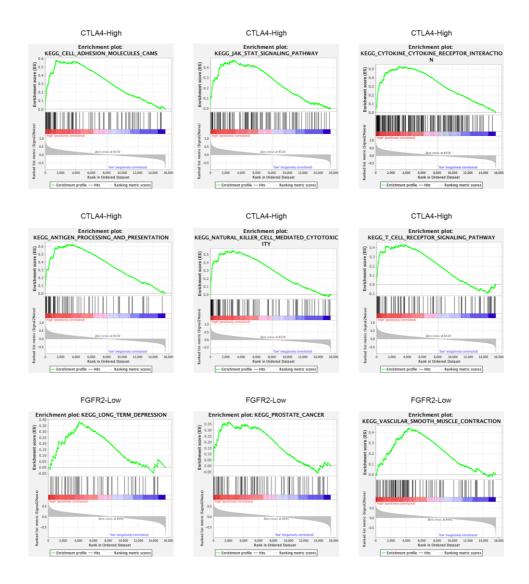


Figure S3. Gene set enrichment analysis (GSEA) for expression profiles in HNSCC samples with relative higher expression of CTLA4 or FGFR2.